

## DATA REPOSITORY

### U/Pb Analytical Procedures

U-Pb TIMS analytical methods utilized in this study are outlined in Parrish et al. (1987). Heavy mineral concentrates were prepared using standard crushing, grinding, Wilfley™ table, and heavy liquid techniques. Mineral separates were sorted by magnetic susceptibility using a Frantz™ isodynamic separator. Single and multigrain zircon fractions analyzed were very strongly air abraded following the method of Krogh (1982). Treatment of analytical errors follows Roddick (1987) with errors on the ages reported at the  $2\sigma$  level (Table 2). U-Pb TIMS concordia diagrams are presented in Fig. 7.

SHRIMP II analyses were conducted at the Geological Survey of Canada (GSC) using analytical procedures described by Stern (1997), with standards and U-Pb calibration methods following Stern and Amelin (2003). Zircons from the samples were cast in 2.5 cm diameter epoxy mounts (GSC mount #284 for samples VLA02178 (z7518) and VL02294 (z7522) and GSC mount #254 for sample RAX01-908 (z7097)) along with fragments of the GSC laboratory standard zircon (z6266, with  $^{206}\text{Pb}/^{238}\text{U}$  age = 559 Ma). The mid-sections of the zircons were exposed using 9, 6, and 1  $\mu\text{m}$  diamond compound, and the internal features of the zircons were characterized with backscatter electrons (BSE) utilizing a Cambridge Instruments scanning electron microscope (SEM). Mount surfaces were evaporatively coated with 10 nm of high purity Au. Analyses were conducted using an  $^{16}\text{O}^-$  primary beam, projected onto the zircons at 10 kV. The sputtered area used for analysis was ca. 25  $\mu\text{m}$  in diameter with a beam current of ca. 6 nA for z7518 and z7522 and ca. 13 nA for z7097. The count rates of ten isotopes of  $\text{Zr}^+$ ,  $\text{U}^+$ ,  $\text{Th}^+$ , and  $\text{Pb}^+$  in zircon were sequentially measured over 6 scans with a single electron multiplier and a pulse counting system with dead time of 35 ns. Off-line data processing was accomplished using customized in-house software. The  $1\sigma$  external errors of  $^{206}\text{Pb}/^{238}\text{U}$  ratios reported in Table 1 incorporate a  $\pm 1.0\%$  error in calibrating the standard zircon (see Stern and Amelin, 2003). No fractionation correction was applied to the Pb-isotope data;

common Pb correction utilized the measured  $^{204}\text{Pb}/^{206}\text{Pb}$  and compositions modeled after Cumming and Richards (1975). The  $^{206}\text{Pb}/^{238}\text{U}$  ages for the analyses have been corrected for common Pb using both the 204- and 207-methods (Stern, 1997), but there is generally no significant difference in the results (Table 2). Isoplot v. 2.49 (Ludwig, 2001) was used to generate concordia plots and calculate weighted means. The data are plotted in concordia diagrams with errors at the  $2\sigma$  level (Fig. 7). A Concordia age (Ludwig, 1998) is calculated for some of the samples presented in this paper. A Concordia age incorporates errors on the decay constants and includes both an evaluation of concordance and an evaluation of equivalence of the data (how well the data fit the assumption that they are repeated measurements of the same point). The calculated Concordia ages and errors quoted in the text are at  $2\sigma$  with decay constant errors included.

### **Sm/Nd Analytical Procedures**

Samples VL01A057, RAX01047, RAX01049, RAX01051, and RAX01059 were analyzed at Geospec (Edmonton Canada). The isotopic composition of Nd was determined in multi-dynamic mode by Thermal Ionization Mass Spectrometry (TIMS). The  $^{143}\text{Nd}/^{144}\text{Nd}$  ratio of samples, normalized for variable mass fractionation, is presented relative to a value of 0.512107 for the Geological Survey of Japan Nd isotopic standard JNd-1 (Tanaka et al., 2000), which is equivalent to a value of 0.511850 for the La Jolla Nd isotopic standard. Sm isotopic abundances were measured in static mode by TIMS and were normalized for variable mass fractionation. For Nd, the long-term average value for JNd-1 is  $0.512110 \pm 0.000012$  (1sd), equivalent to a value of 0.511853 for the La Jolla Nd isotopic standard, and within the generally accepted window of La Jolla values ( $0.511850 \pm 0.000010$ ). For Sm, the long-term average value for  $^{149}\text{Sm}/^{154}\text{Sm}$  determined from a Sm standard solution was  $0.60759 \pm 0.00009$  (1sd), identical within quoted uncertainty to the value determined by Wasserburg et al. (1981) of  $0.60750 \pm 0.00002$ .

Samples VL01A097a, VL01A097a, VL01A360, VL02A283, VL02A295 were analyzed at Ottawa-Carleton Geoscience Centre Radiogenic Isotope Research Laboratory (Ottawa, Canada) using a Finnigan MAT261 multicollector mass spectrometer equipped with an electron multiplier. Samples were spiked with a mixed  $^{148}\text{Nd}$ - $^{149}\text{Sm}$  spike prior to dissolution. Concentrations were precise to + 1%, but  $^{147}\text{Sm}/^{144}\text{Nd}$  ratios were reproducible to 0.5%. Samples were loaded with 0.25N HCl on one side of a Re double filament, and run at temperatures of 1780-1820°C. Isotope ratios were normalized to  $^{146}\text{Nd}/^{144}\text{Nd} = 0.72190$ . Analyses of the USGS standard BCR-1 yield Nd = 29.02 ppm, Sm = 6.68 ppm, and  $^{143}\text{Nd}/^{144}\text{Nd} = 0.512668 +/- 20$  (n=4). 32 runs of the La Jolla standard average  $^{143}\text{Nd}/^{144}\text{Nd} = 0.511875 +/- 18$  (Sept. 1992-Sept. 1994).

### **Geochemistry Analytical Procedures**

Multiple laboratories, listed in Table DR1, were employed for analysis of samples. Detailed information on sample preparation and analytical procedures is available in Rogers (2004). Detection limits for all laboratories are listed in Table DR1. McGill University Geochemical Laboratories (Montreal, Canada) were utilized for XRF determination of major and select trace elements on glass pellets using PHILIPS PW2440 4kW automated XRF spectrometer, with accuracy within 1%. Ontario Geological Survey Laboratories (Sudbury, Canada) determined major and select trace elements on the Rigaku RIX-3000 WD-XRF; trace elements were analyzed on the Perkin-Elmer ELAN 5000 ICP-MS. Precision and accuracy are described in Richardson et al. (1996). Geological Survey of Canada Analytical Chemistry Laboratory (Ottawa, Canada) was utilized for both XRF (major elements) and ICP-MS (trace elements). The precision for trace elements is generally better than 10%. X-Ray Laboratories (XRAL, Toronto, Canada) determined major elements in duplicate on fused glass discs. Pressed powder pellets were utilized for select trace elements. Analysis of trace elements at Memorial University of Newfoundland (St. John's, Canada) followed the methods and had accuracy and precision outlined in Jenner et al. (1990).

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Table DR1: Geochemistry of peri-Laurentian arc - back-arc complexes in central Newfoundland

Sample	RAX00 705	RAX02 169	VL01 076	VL01 079	VL01A 071	VL01A 058	VL01A 277	VL01A 207b	VL01A 122a	VL01A 351	VL01A 352	VL01A 264	VL01A 057
UTMx <sup>†</sup>	440769	483605	442300	443650	449441	460378	461124	453940	447878	467183	467119	459504	460403
UTMy <sup>†</sup>	5336018	5376892	5337361	5338411	5342789	5350044	5350959	5345097	5342647	5363419	5363458	5349121	5349812
NTS map Unit	12A/04 SB <sub>1</sub>	12A/11 SB <sub>1</sub>	12A/04 SB <sub>1</sub>	12A/04 SB <sub>1</sub>	12A/05 SB <sub>1</sub>	12A/05 SB <sub>1</sub>	12A/05 SB <sub>1</sub>	12A/04 SB <sub>1</sub>	12A/04 SB <sub>1</sub>	12A/06 SB <sub>1</sub>	12A/06 SB <sub>1</sub>	12A/05 SB <sub>1</sub>	12A/05 SB <sub>1</sub>
Rock type <sup>‡</sup>	pbslt	bslt	diab	gabb	diab	bslt	bslt	bslt	bslt	gabb	diab	bslt	diab
Laboratory <sup>§</sup>	1	4	5	5	5	5	5	5	5	6	3	5	5
SiO <sub>2</sub> (wt%)	48.70	47.96	48.92	50.80	49.50	47.83	49.64	49.99	46.09	45.88	47.78	47.28	46.98
TiO <sub>2</sub>	1.22	1.57	1.08	1.46	1.30	1.37	1.40	1.27	1.35	1.42	1.11	1.71	2.00
Al <sub>2</sub> O <sub>3</sub>	20.20	14.38	16.34	16.36	15.60	14.77	13.95	14.22	17.60	14.23	16.04	14.68	14.26
Fe <sub>2</sub> O <sub>3</sub> total	10.40	12.81	9.46	10.98	9.54	11.56	11.56	11.46	10.95	11.47	11.21	12.79	14.92
MnO	0.20	0.19	0.17	0.18	0.21	0.27	0.18	0.24	0.18	0.18	0.23	0.22	0.23
MgO	4.54	7.49	7.50	5.13	7.04	8.22	7.01	8.61	8.54	8.09	8.19	7.70	8.83
CaO	3.99	10.76	12.34	6.71	10.64	9.84	12.10	7.59	9.05	13.08	10.23	9.79	6.24
Na <sub>2</sub> O	4.01	2.95	2.44	6.08	4.05	3.73	2.59	4.73	3.18	1.72	2.98	3.60	3.74
K <sub>2</sub> O	2.15	0.06	0.33	0.14	0.12	0.05	0.08	0.04	0.28	0.57	0.49	0.08	0.08
P <sub>2</sub> O <sub>5</sub>	0.08	0.13	0.08	0.10	0.10	0.10	0.11	0.10	0.10	0.12	0.10	0.15	0.14
LOI	6.71	2.43	2.48	2.66	2.57	2.97	2.24	2.94	3.82	3.00	2.09	3.05	3.90
Total	99.00	100.81	101.12	100.60	100.66	100.70	100.85	101.17	101.15	99.81	100.53	101.04	101.32
Ba (ppm)	202	15.1	53.6	78.6	57.0	12.9	25.1	25.7	57.1	43.0	33.2	26.1	40.2
Cr	397	271	271	265	274	287	213	69	286	226	347	235	130
Cs	2.3	0.18	0.66	0.37	0.13	0.86	0.44	0.38	0.29	3.4	2.8	0.96	0.20
Hf	1.60	2.70	1.80	2.49	2.39	2.62	2.41	2.00	2.19	3.00	1.70	3.11	3.40
Nb	2.40	2.14	2.52	3.28	3.48	2.83	3.11	3.40	3.74	5.40	2.98	5.32	3.44
Ni	146	88	97	104	95	102	64	56	110	46	105	90	61
Pb	b.d.	1	13.46	4.15	2.52	0.25	0.36	1.68	0.97	1.3	6.3	0.36	0.37
Rb	50.0	0.47	9.55	2.9	1.49	0.82	1.2	0.44	6.44	35.2	25.7	0.91	1.4
Sc	41	47.2	43.4	42.9	39.1	43.5	39.4	43.6	43.2	43	52.5	42.4	48.7
Sr	143	140	134	209	401	119	184	137	166	196	204	130	90.9
Ta	0.23	0.26	b.d.	0.4	0.39	0.3	b.d.						
Th	0.14	0.17	0.18	0.22	0.24	0.23	0.22	0.22	0.27	0.3	0.23	0.39	0.25
U	0.11	0.06	0.06	0.08	0.08	0.07	0.07	0.08	0.08	b.d.	0.07	0.12	0.08
V	186	346	251	314	255	294	295	296	300	313	315	345	351
Y	35	41	24.86	31.06	29.79	34.1	32.76	29.32	30.75	32.2	26.6	39.84	47.58
Zr	52	91.5	58.67	81.06	84.69	87.48	82.35	69.74	77.32	74.6	65.7	112.97	119.59
La	3.80	3.02	2.70	3.21	3.58	2.96	3.14	2.92	3.42	4.90	2.90	4.97	3.79
Ce	9.60	9.60	7.40	9.55	10.46	9.20	9.51	8.46	9.79	12.70	7.97	13.96	11.83
Pr	1.90	1.81	1.31	1.64	1.82	1.69	1.70	1.49	1.66	2.04	1.41	2.38	2.19
Nd	10.00	10.36	7.32	9.03	9.64	9.20	8.89	7.93	8.63	9.80	7.59	12.18	12.07
Sm	3.40	3.72	2.52	3.14	3.32	3.48	3.26	2.89	2.96	3.40	2.58	4.13	4.60
Eu	1.30	1.44	1.10	1.32	1.42	1.21	1.28	1.15	1.24	1.10	0.99	1.49	1.52
Gd	4.90	5.95	3.84	4.68	4.50	4.90	4.55	3.96	4.24	4.48	3.71	5.64	6.51
Tb	0.94	1.04	0.66	0.82	0.77	0.88	0.86	0.74	0.82	0.75	0.67	1.04	1.17
Dy	5.70	6.89	4.36	5.55	5.00	5.95	5.66	4.84	5.22	5.37	4.51	6.77	7.79
Ho	1.20	1.54	0.99	1.22	1.10	1.29	1.29	1.10	1.19	1.09	0.97	1.56	1.73
Er	3.10	4.47	2.88	3.52	3.14	3.62	3.60	3.05	3.36	3.08	2.83	4.35	5.12
Tm	0.50	0.66	0.43	0.52	0.45	0.57	0.55	0.48	0.52	0.56	0.41	0.67	0.74
Yb	3.20	4.11	2.77	3.33	2.92	3.58	3.50	2.93	3.20	3.26	2.72	4.16	4.75
Lu	0.44	0.66	0.42	0.50	0.44	0.56	0.55	0.47	0.52	0.52	0.42	0.66	0.73
Mg# <sup>††</sup>	48.8	56.0	63.3	50.5	61.7	60.8	56.9	62.1	63.0	60.6	61.4	56.8	56.3
(La/Th)n <sup>#</sup>	1.3	0.9	0.7	0.7	0.7	0.6	0.7	0.6	0.6	0.8	0.6	0.6	0.7
(La/Nb)n <sup>#</sup>	1.5	1.3	1.0	0.9	1.0	1.0	0.9	0.8	0.9	0.8	0.9	0.9	1.0
(Zr/Sm)n <sup>#</sup>	0.54	0.87	0.83	0.92	0.91	0.89	0.90	0.86	0.93	0.78	0.91	0.97	0.92
(La/Yb)n <sup>#</sup>	1.4	0.9	1.2	1.2	1.5	1.0	1.1	1.2	1.3	1.8	1.3	1.5	1.0

† UTM zone 21 (NAD83); ‡ (p)bslt-(pillowed)basalt; (s)diab-(sheeted)diabase; gabb-gabbro; (f,l,m)tuff-(felsic,intermediate,mafic)tuff; gran-granodiorite; rhyo-rhyolite; fdyke-felsic dyke; (p)and-(pillowed)andesite; § Laboratory 1:GSC(2003); 2:OGS(2003); 3:McGill; OGS (2002); 4:McGill; OGS (2001); 5:OGS (2001)

6:Acme (2001); 7:XRAL Toronto; MUN (1992); 1 to 6: Rogers (2004); 7: Jenner et al., (1990); # b.d. - below detection; n.a. not applicable; †† Mg# =

Mg<sup>2+</sup>/(Mg<sup>2+</sup>+Fe<sup>2+</sup>), Fe<sup>3+</sup>/Fe<sup>2+</sup>=0.1; ‡‡ N-MORB normalized value (Sun and MacDonough, 1989)

Table DR1 (cont'd)

VL01A 271	VL01A 265	VL01J 164b	VL02A 211	VL02A 005	VL02A 292	VL02A 232b	VL02A 171	VL02A 276	VL02A 237	VL02A 222b	VL02A 285	VL02A 214	VL02A 275a	VL02A 298b
459998	459501	467178	483594	439860	482796	475530	475002	483838	471437	477731	480662	483699	484076	484922
5349786	5349209	5361549	5376878	5335742	5376260	5370224	5369441	5376862	5367487	5371616	5373460	5376546	5377009	5378280
12A/05	12A/05	12A/06	12A/11	12A/04	12A/11	12A/06	12A/06	12A/11	12A/06	12A/06	12A/11	12A/11	12A/11	12A/11
SB <sub>1</sub>														
bslt	bslt	diab	bslt	diab	bslt	diab	gabb	bslt	bslt	gabb	bslt	bslt	diab	bslt
1	1	5	3	3	3	3	3	3	3	3	3	3	3	3
47.80	47.60	47.25	48.13	47.57	48.34	49.45	49.61	46.54	49.04	50.16	49.80	48.70	48.12	48.13
1.73	2.00	1.39	0.98	1.26	1.39	1.51	1.25	1.27	1.41	1.58	1.58	1.51	1.48	1.68
13.70	13.90	15.30	15.63	15.26	14.62	16.09	14.92	15.70	13.83	13.46	13.84	14.22	14.16	14.26
13.30	14.00	11.25	10.11	11.23	11.54	11.24	10.71	10.94	11.91	12.70	12.66	12.75	12.43	13.53
0.20	0.20	0.21	0.19	0.16	0.18	0.20	0.21	0.17	0.25	0.20	0.20	0.21	0.20	0.22
7.63	7.85	7.95	8.90	8.68	7.43	6.44	8.39	7.30	7.33	7.30	7.41	7.56	7.89	7.73
8.90	7.66	12.16	11.23	10.15	11.35	9.77	10.18	11.69	10.48	10.22	9.29	10.13	10.69	8.01
3.34	3.52	2.39	2.80	3.03	3.06	4.19	3.35	3.06	3.99	3.02	3.62	3.36	3.00	3.44
0.10	0.07	0.07	0.12	0.10	0.14	0.11	0.10	0.06	0.13	0.11	0.21	0.06	0.09	0.13
0.13	0.15	0.11	0.09	0.11	0.12	0.15	0.11	0.12	0.12	0.14	0.13	0.12	0.14	0.14
6.01	6.96	2.15	2.37	3.05	2.23	1.73	1.42	3.58	1.81	1.94	1.79	2.15	2.29	3.36
99.30	99.90	100.22	100.63	100.67	100.48	100.96	100.33	100.52	100.33	100.90	100.59	100.83	100.58	100.69
31.0	44.0	11.5	24.8	15.4	10.4	16.7	18.4	9.8	31.8	19.1	15.3	18.4	13.1	27.7
150	160	278	399	309	309	420	357	69	188	178	213	364	209	
0.45	0.32	0.44	0.20	0.78	0.18	0.11	0.22	0.14	0.11	0.19	0.50	0.14	0.13	0.15
2.60	3.20	2.55	1.60	2.00	2.30	2.60	2.20	2.20	2.30	2.60	2.70	2.50	2.40	2.70
3.30	4.10	3.28	2.42	2.22	2.65	4.41	2.62	4.72	3.55	3.19	2.43	2.34	4.94	2.49
71	73	94	169	112	94	118	124	122	51	76	74	78	107	77
0.5	0.5	0.83	0.7	6.4	1.5	1.3	0.5	1.4	0.8	1.6	2	1.3	1	0.8
1.3	0.72	1.44	1.12	1.68	0.77	0.5	1.38	0.5	0.44	1.05	4.39	0.23	0.18	1.46
44	44	43.1	39.6	43.7	43.5	42.5	47.8	43.9	46.2	43.9	48.9	49.6	44.2	45.5
90	157	173	167	104	131	151	163	220	269	183	177	200	217	129
0.21	0.25	b.d.	0.27	0.18	0.3	0.4	0.28	0.45	0.37	0.33	0.31	0.3	0.41	0.31
0.19	0.25	0.22	0.19	0.18	0.18	0.31	0.18	0.33	0.26	0.24	0.2	0.19	0.34	0.18
0.07	0.08	0.07	0.05	0.05	0.07	0.26	0.06	0.25	0.09	0.09	0.07	0.07	0.11	0.06
357	385	293	254	294	304	307	315	315	315	334	337	343	344	356
43	50	32.05	22.32	29.79	34.8	34.79	30.38	31.49	33.52	38.31	38.42	37.95	38.08	41.01
91	105	85.81	56.2	76.2	77.2	93.2	76.1	76.4	86	97.5	91	91	92	95.2
3.60	4.50	3.35	2.48	2.74	2.79	4.46	3.03	4.16	3.36	3.56	3.12	2.92	4.67	3.14
11.00	14.00	10.02	7.23	8.19	8.52	12.50	9.13	10.95	10.22	10.75	9.70	9.38	12.37	10.04
1.90	2.40	1.74	1.19	1.44	1.55	2.08	1.55	1.77	1.77	1.92	1.85	1.67	1.96	1.88
11.00	14.00	10.03	6.58	8.18	8.78	10.81	8.95	9.58	9.82	10.61	10.25	9.80	11.33	10.50
3.70	4.60	3.51	2.37	2.80	3.30	3.49	3.12	3.19	3.35	3.81	3.74	3.48	3.56	3.80
1.20	1.40	1.39	0.95	0.98	1.23	1.46	1.15	1.21	1.15	1.40	1.26	1.23	1.31	1.40
5.60	6.70	4.66	3.35	4.30	4.77	5.13	4.38	4.47	4.80	5.28	5.28	5.42	5.41	5.77
1.10	1.20	0.85	0.58	0.76	0.93	0.96	0.83	0.85	0.83	1.00	0.95	0.99	0.92	1.10
6.70	7.80	5.48	3.92	4.91	6.03	5.81	5.34	5.39	5.81	6.65	6.43	6.44	6.14	6.95
1.50	1.70	1.24	0.90	1.09	1.31	1.27	1.15	1.19	1.21	1.46	1.43	1.42	1.40	1.51
4.00	4.70	3.58	2.57	3.23	3.74	3.86	3.29	3.39	3.84	4.35	4.24	4.39	4.26	4.64
0.65	0.75	0.55	0.37	0.48	0.59	0.56	0.49	0.53	0.55	0.63	0.61	0.64	0.64	0.68
4.20	5.00	3.41	2.41	3.13	3.67	3.40	3.08	3.30	3.43	3.98	3.82	3.77	3.95	4.26
0.69	0.77	0.51	0.37	0.49	0.55	0.52	0.49	0.49	0.52	0.64	0.61	0.59	0.62	0.64
55.6	55.0	60.6	65.7	62.8	58.4	55.5	63.1	59.3	57.3	55.6	56.1	56.4	58.0	55.5
0.9	0.9	0.7	0.6	0.7	0.7	0.7	0.8	0.6	0.6	0.7	0.7	0.7	0.7	0.8
1.0	1.0	1.0	1.0	1.2	1.0	0.9	1.1	0.8	0.9	1.0	1.2	1.2	0.9	1.2
0.87	0.81	0.87	0.84	0.97	0.83	0.95	0.87	0.85	0.91	0.91	0.86	0.93	0.92	0.89
1.0	1.1	1.2	1.3	1.1	0.9	1.6	1.2	1.5	1.2	1.1	1.0	0.9	1.4	0.9

Table DR1 (cont'd)

VL02A 297	VL02A 291	VL01A 272	VL01J 107	VL02A 287	RAX02 170	VL01A 59	VL01J 063a	VL02A 225	VL02A 223b	VL02A 233a	VL02A 206	VL02J 404	VL01A 283a	VL01J 151
484863	482863	459932	466735	481406	483690	460416	448841	477142	477578	475514	484714	495564	465013	474143
5378393	5375733	5349869	5362789	5374122	5376649	5350068	5344158	5371257	5371723	5370286	5378095	5398809	5357399	5370148
12A/11	12A/11	12A/05	12A/06	12A/11	12A/11	12A/05	12A/04	12A/06	12A/06	12A/06	12A/11	12A/11	12A/06	12A/06
SB <sub>1</sub>	SB <sub>1</sub>	SB <sub>2</sub>	SB <sub>2</sub>	SB <sub>2</sub>	SB <sub>3</sub>	OBS	OBS							
diab	bslt	bslt	diab	sdiab	bslt	diab	diab	gabb	gabb	bslt	diab	bslt	bslt	diab
3	3	6	5	3	4	1	5	3	3	3	3	3	5	5
46.56	46.83	48.14	48.26	47.17	48.49	45.80	47.45	48.18	46.96	48.09	47.90	47.04	49.33	49.68
1.70	1.91	1.33	1.72	1.59	1.00	0.54	0.75	0.38	0.44	1.01	1.30	0.72	1.51	1.30
14.53	14.47	15.07	14.53	14.29	15.04	15.00	16.83	16.58	14.03	15.95	14.80	17.06	14.52	15.54
13.65	14.76	10.93	12.01	12.67	11.94	10.10	9.80	7.67	10.64	10.33	11.44	9.25	11.77	12.02
0.21	0.25	0.18	0.22	0.19	0.16	0.16	0.16	0.14	0.19	0.15	0.19	0.18	0.20	0.19
8.76	7.62	7.07	7.67	7.63	8.07	11.10	8.99	10.46	11.85	6.05	7.93	9.20	6.16	8.24
8.55	9.36	12.04	11.65	12.57	8.80	10.50	12.90	12.28	12.37	10.87	11.44	11.65	12.23	8.62
2.96	3.10	2.32	2.46	2.03	2.58	1.66	1.80	2.13	1.36	3.62	2.63	1.82	3.05	4.28
0.08	0.04	0.06	0.34	0.08	1.04	0.49	0.02	0.24	0.11	0.12	0.10	0.97	0.05	0.10
0.14	0.16	0.16	0.11	0.13	0.06	0.01	0.04	0.02	0.03	0.07	0.11	0.04	0.17	0.14
3.24	2.63	2.50	1.28	2.14	3.43	9.25	1.42	2.42	2.59	3.92	2.43	2.19	1.59	0.58
100.45	101.19	99.86	100.26	100.56	100.70	100.00	100.16	100.59	100.64	100.26	100.34	100.24	100.57	100.70
25.6	10.7	23.0	21.7	11.9	45.7	73.0	b.d.	14.0	9.4	12.4	19.2	156	10.0	6.2
211	152	267	216	284	420	194	434	410	284	368	360	524	174	253
0.19	0.12	0.30	2.1	0.17	0.38	0.62	0.23	0.63	0.19	0.13	0.20	3.8	0.03	0.09
2.90	3.40	2.40	1.46	1.80	1.40	0.37	0.99	0.40	0.70	1.60	2.10	0.90	2.44	2.52
2.44	3.23	2.70	4.02	1.89	0.97	0.33	0.47	0.20	0.36	0.83	1.70	0.18	4.81	1.79
73	72	39	82	89	99	79	157	167	161	83	95	211	87	91
b.d.	b.d.	0.7	3.15	1.3	b.d.	0.5	b.d.	1.1	1.1	1	0.5	6.3	2.79	3.49
1.25	0.24	0.25	23.2	0.52	20.4	14.0	0.28	8.82	1.09	1.59	0.8	42.8	b.d.	0.75
49.1	52.5	35	46.8	43.8	48.3	53	37.1	40.8	48.8	41.6	52.5	32.8	37.7	40.5
73.9	81.4	126	174	119	135	82	68.3	123	84.5	172	147	72.4	291	92.3
0.28	0.33	0.2	b.d.	0.27	0.19	0.025	b.d.	0.14	0.14	0.19	0.23	b.d.	b.d.	b.d.
0.2	0.23	b.d.	b.d.	b.d.	0.08	0.03	b.d.	b.d.	b.d.	0.07	0.13	b.d.	0.79	1.08
0.06	0.10	b.d.	0.02	0.01	0.08	b.d.	0.01	0.01	0.02	0.03	0.04	b.d.	0.3	0.32
361	404	272	351	330	289	320	219	166	202	263	315	209	331	313
42.96	47.67	29.3	29.7	34	23.74	7.3	20.23	10.66	16.08	25.71	32.56	20.07	30.78	33.54
100.2	114.5	73.5	43.67	49.6	45.1	11	28.89	9.8	20.2	53.4	81.9	22.3	89.66	94.29
3.32	4.03	4.60	2.64	2.27	1.28	0.30	0.77	0.38	0.72	1.29	2.47	0.33	7.86	6.32
10.49	12.23	11.80	7.93	7.25	4.13	1.00	2.63	1.33	2.42	4.83	8.12	1.49	18.88	16.29
1.94	2.25	1.89	1.41	1.45	0.77	0.22	0.54	0.30	0.52	0.99	1.55	0.38	2.81	2.49
10.98	12.72	10.90	8.20	8.68	4.66	1.20	3.40	1.94	3.12	6.20	8.77	2.92	13.22	12.32
4.08	4.71	3.30	3.14	3.27	1.94	0.55	1.61	0.84	1.19	2.42	3.33	1.54	3.84	3.73
1.43	1.45	1.13	1.38	1.36	0.85	0.22	0.72	0.47	0.58	0.97	1.23	0.60	1.63	1.26
6.17	6.58	4.27	4.53	4.99	3.14	0.98	2.53	1.40	2.04	3.60	4.38	2.60	4.82	4.81
1.10	1.22	0.81	0.80	0.93	0.58	0.19	0.47	0.30	0.42	0.68	0.80	0.50	0.86	0.84
7.32	8.22	5.08	5.17	5.89	3.80	1.30	3.28	1.94	2.60	4.25	5.52	3.40	5.45	5.53
1.61	1.79	1.03	1.10	1.29	0.90	0.28	0.76	0.41	0.58	0.94	1.18	0.76	1.21	1.22
4.78	5.36	2.75	3.15	3.79	2.64	0.82	2.17	1.29	1.88	2.67	3.45	2.25	3.27	3.54
0.71	0.80	0.49	0.46	0.55	0.42	0.12	0.33	0.17	0.25	0.40	0.53	0.33	0.50	0.53
4.34	4.94	2.80	2.76	3.60	2.52	0.80	2.12	1.15	1.70	2.64	3.01	2.00	3.10	3.44
0.68	0.78	0.53	0.41	0.56	0.42	0.12	0.31	0.18	0.27	0.39	0.50	0.33	0.48	0.54
58.3	52.9	58.5	58.2	56.8	59.6	70.5	66.7	74.8	70.8	56.1	60.2	68.4	53.3	59.9
0.8	0.8	n.a. <sup>#</sup>	n.a.	n.a.	0.8	0.5	n.a.	n.a.	n.a.	0.9	0.9	n.a.	0.5	0.3
1.3	1.2	1.6	0.6	1.1	1.2	0.8	1.5	1.8	1.9	1.4	1.4	1.7	1.5	3.3
0.87	0.86	0.79	0.49	0.54	0.83	0.71	0.64	0.41	0.60	0.78	0.87	0.51	0.83	0.90
0.9	1.0	2.0	1.2	0.8	0.6	0.5	0.4	0.4	0.5	0.6	1.0	0.2	3.1	2.2

Table DR1 (cont'd)

VL01J 208a	VL01J 208b	VL02A 277	VL02A 255	VL02A 192a	VL02J 402	VL02J 403	WXNF 173	VL01A 097b	VL01 050	VL01 083	VL01 082	VL01A 099	VL01A 098b	VL01A 097a
495317	495317	483312	487807	474119	494496	494805	494350	436177	436019	435770	440850	435221	435139	436177
5397037	5397037	5376757	5383738	5369669	5398217	5398458	5396260	5332496	5333068	5332411	5337211	5331405	5331286	5332496
12A/11	12A/11	12A/11	12A/11	12A/06	12A/11	12A/11	12A/11	12A/04						
OBS	OBS	OBS	OBS	OBS	OBS	OBS	OBS	ML <sub>1</sub>	ML <sub>2</sub>					
bslt	bslt	sdiab	diab	diab	bslt	bslt	ftuff	pbslt	pbslt	pbslt	pbslt	pbslt	pbslt	bslt
5	5	3	3	3	3	3	7	5	5	5	5	3	3	3
48.24	47.79	49.01	49.09	47.11	47.59	47.52	48.2	72.76	54.95	50.27	47.12	49.57	51.95	52.78
3.07	3.10	2.36	1.50	2.16	2.42	2.93	1.59	0.11	0.65	0.61	0.55	0.63	0.70	0.71
13.48	12.91	14.70	15.04	15.34	14.08	13.05	14.65	11.83	14.97	16.94	15.34	14.71	14.66	16.48
16.41	18.45	12.82	13.16	14.48	13.62	16.11	12.05	4.25	9.83	9.97	9.80	9.83	9.72	11.77
0.23	0.22	0.22	0.17	0.18	0.28	0.30	0.21	0.05	0.12	0.16	0.20	0.16	0.20	0.20
5.59	4.53	5.77	6.68	4.94	7.27	5.91	6.32	1.70	5.06	7.13	7.50	5.97	5.99	4.94
5.45	5.70	10.02	11.82	11.88	9.72	8.51	8.81	4.10	8.00	6.88	9.03	6.33	6.35	6.90
5.31	5.46	3.51	2.41	2.95	3.60	4.36	3.15	0.43	2.92	4.98	2.53	1.15	4.44	2.37
0.05	0.06	0.29	0.11	0.18	0.36	0.21	0.34	2.84	0.04	0.24	1.05	2.87	0.07	1.33
0.41	0.40	0.27	0.20	0.27	0.26	0.32	0.2	0.02	0.07	0.08	0.07	0.06	0.10	0.10
2.48	2.14	2.01	0.47	0.40	0.98	1.32	3.1	2.56	4.11	4.07	6.98	8.89	6.27	2.73
100.71	100.75	101.03	100.73	99.94	100.25	100.58	98.62	100.64	100.71	101.32	100.18	100.20	100.48	100.35
10.6	8.4	165	31.6	16.5	61.9	21.4	113	147	14.0	31.9	209	207	37.4	290
11	14	114	214	56	200	55	190	3	26	89	56	50	32	37
0.13	0.12	0.29	0.12	0.24	0.92	0.19	b.d.	2.9	0.28	0.13	0.68	3.8	0.62	1.9
5.19	4.60	5.00	2.20	5.00	4.50	5.10	3.01	3.74	1.04	1.05	0.87	1.10	1.30	1.20
7.95	7.21	6.22	5.23	2.44	4.47	7.80	6.90	3.85	0.66	0.59	0.64	0.66	0.83	0.89
15	16	32	67	35	58	39	43	b.d.	22	43	37	30	19	19
0.69	0.48	4.1	b.d. <sup>#</sup>	2	4.1	4.4	b.d.	12.02	0.67	0.25	4.18	4.3	4.9	1.6
0.21	b.d.	3.57	0.72	1.98	5.96	2.22	5	72.9	0.73	2.44	20.7	71.2	0.68	21.3
42.9	43.2	44.6	44.2	43.4	45.5	38.2	33	18.9	40.2	43.8	44.6	49.6	45.1	52.5
32.8	35.5	254	178	187	240	103	201	444	71.1	85.0	233	114	468	227
0.48	0.47	0.57	0.41	0.28	0.3	0.59	b.d.	b.d.	b.d.	b.d.	b.d.	0.22	0.24	0.24
0.92	0.8	1.82	0.85	0.93	0.44	0.66	0.53	3.66	0.76	0.56	0.8	0.74	0.74	1.05
0.42	0.37	0.68	0.27	0.46	0.22	0.3	b.d.	0.83	0.23	0.53	0.33	0.13	0.22	0.27
351	351	315	367	417	388	446	341	14	234	236	294	315	315	324
48.56	41.7	45.6	28.97	56.96	45.2	50.29	27	32.7	16.58	15.36	13.85	14.11	17.58	19.38
210.12	184.16	190.1	81.5	180.1	171.3	196.2	111	119.42	34.43	32.28	24.87	36.8	43.9	44
10.71	8.90	12.64	8.55	7.96	7.13	10.09	6.23	12.12	4.37	3.06	4.41	3.72	4.58	5.65
29.10	24.67	30.97	19.84	22.80	21.23	28.39	15.76	30.82	10.02	7.40	9.89	8.53	10.45	12.32
4.43	3.93	4.78	2.88	3.79	3.60	4.49	2.48	4.23	1.45	1.10	1.44	1.27	1.53	1.73
22.72	20.48	22.90	13.47	20.38	18.63	22.69	12.4	18.01	6.77	5.41	6.67	5.73	7.26	7.84
6.45	5.97	6.66	3.64	6.29	5.94	6.72	3.77	4.66	1.97	1.76	1.94	1.79	2.14	2.18
2.35	2.25	2.06	1.25	2.28	1.94	2.41	1.36	0.91	0.72	0.65	0.76	0.64	0.84	0.70
8.05	6.97	7.70	4.60	8.50	7.79	8.71	4.58	5.20	2.60	2.31	2.23	2.15	2.64	2.78
1.34	1.19	1.33	0.79	1.56	1.26	1.45	0.78	0.86	0.46	0.42	0.36	0.38	0.46	0.46
8.16	7.58	8.30	5.28	9.70	7.98	9.05	5.25	5.24	2.82	2.72	2.40	2.44	2.93	3.00
1.75	1.58	1.72	1.09	2.19	1.72	1.94	1.1	1.17	0.63	0.58	0.54	0.54	0.67	0.68
4.92	4.63	4.91	3.23	6.23	5.00	5.59	3.13	3.45	1.82	1.78	1.61	1.70	2.00	1.92
0.72	0.67	0.69	0.46	0.98	0.73	0.83	0.45	0.56	0.29	0.28	0.24	0.25	0.32	0.31
4.44	4.06	4.40	2.94	5.93	4.38	5.05	2.97	3.85	1.88	1.78	1.56	1.53	1.92	2.07
0.68	0.63	0.66	0.45	0.90	0.68	0.82	0.44	0.64	0.28	0.28	0.24	0.25	0.31	0.33
42.6	34.9	49.5	52.5	42.6	53.8	44.4	53.4	46.6	52.9	60.9	62.5	57.0	57.3	47.8
0.6	0.5	0.3	0.5	0.4	0.8	0.7	0.6	0.2	0.3	0.3	0.3	0.2	0.3	0.3
1.3	1.2	1.9	1.5	3.0	1.5	1.2	0.8	2.9	6.2	4.8	6.4	5.3	5.1	5.9
1.16	1.10	1.01	0.80	1.02	1.02	1.04	1.05	0.91	0.62	0.65	0.46	0.73	0.73	0.72
2.9	2.7	3.5	3.5	1.6	2.0	2.4	2.6	3.8	2.8	2.1	3.4	3.0	2.9	3.3

Table DR1 (cont'd)

VL02A 015	VL02A 016a	RAX01 59	RAX02 158	VL02A 293	RAX02 157	RAX02 156	RAX01 100	VL01 085	VL01A 125	VL01A 297	VL01A 303	VL01A 126	VL01A 131b	VL01A 299b
435952	435865	506714	506083	505825	507469	507598	501123	446650	447564	466268	466033	447772	446449	465979
5332145	5332372	5396042	5397449	5396324	5398957	5399109	5383334	5340686	5341725	5359816	5357613	5341414	5340538	5358413
12A/04	12A/04	12A/10	12A/10	12A/10	12A/10	12A/10	12A/10	12A/04	12A/04	12A/06	12A/06	12A/04	12A/04	12A/06
ML <sub>2</sub>	ML <sub>2</sub>	SK <sub>1</sub>	SK <sub>1</sub>	SK <sub>1</sub>	SK <sub>2</sub>	SK <sub>2</sub>	HR <sub>1</sub>							
bslt	pbslt	pbslt	pbslt	bslt	bslt	bslt	tuff	pbslt	bslt	bslt	bslt	pbslt	pbslt	bslt
3	3	2	4	3	6	4	2	5	5	5	3	3	3	3
47.58	49.51	50.90	51.52	46.47	61.88	45.17	54.00	46.70	47.16	51.25	46.22	50.36	51.09	48.57
0.54	0.65	1.08	0.75	0.66	1.16	1.57	1.44	0.80	1.23	1.23	1.40	1.47	1.33	1.68
14.59	17.47	16.06	15.07	15.99	15.83	17.04	13.54	18.46	15.86	15.61	14.97	14.14	15.42	13.87
9.09	12.17	11.73	12.31	9.49	6.31	16.14	11.37	7.21	11.13	9.79	11.68	11.85	10.75	13.41
0.17	0.19	0.14	0.44	0.17	0.10	0.17	0.16	0.14	0.22	0.14	0.20	0.20	0.16	0.18
4.55	4.92	7.25	8.85	9.75	4.23	7.58	6.72	6.34	7.21	6.64	8.08	6.28	5.34	6.90
15.96	7.63	5.22	3.47	12.21	0.74	3.26	5.22	9.61	12.28	8.32	13.29	9.85	7.48	9.86
0.18	2.42	5.47	1.30	1.35	6.50	4.24	6.29	4.52	3.05	4.86	2.42	4.48	5.72	3.47
0.07	2.09	0.07	0.31	1.23	0.04	0.54	0.43	0.03	0.03	0.09	0.04	0.14	0.06	0.13
0.10	0.08	0.11	0.12	0.08	0.36	0.05	0.09	0.05	0.10	0.09	0.12	0.12	0.12	0.14
7.25	3.52	3.25	5.88	2.93	2.70	5.04	2.16	6.40	2.49	2.59	2.10	1.93	3.10	2.19
100.11	100.69	101.27	100.06	100.46	99.86	100.83	101.42	100.27	100.76	100.60	100.61	100.91	100.64	100.46
10.3	329	41.0	27.3	193	44.0	341	26.8	13.5	23.4	22.7	7.8	80.6	43.4	27.9
5	21	21	165	573	3	b.d.	154	371	264	221	420	366	274	178
0.26	1.0	0.04	0.38	0.41	b.d.	0.19	0.23	0.15	0.19	0.24	0.14	0.66	0.17	0.31
0.90	0.90	1.50	1.20	1.10	2.70	2.00	2.41	1.24	2.15	2.02	2.30	2.50	2.30	2.90
0.66	1.29	0.92	1.16	0.70	4.80	3.72	1.98	1.08	3.87	2.98	2.23	2.42	3.00	2.66
27	3	21	48	196	2	3	67	150	90	57	128	121	97	74
2.2	1.2	1.03	1.9	1.2	2.6	5	0.44	0.56	1.11	0.46	2	0.5	1.2	1.2
1.28	39.5	0.68	4.81	17.1	1.2	3.19	6.72	0.48	0.26	1.2	0.31	2.43	0.41	2.29
37	39.5	38.3	42.3	42.0	23	45.5	46.5	29.0	41.3	41.4	52.5	52.5	52.5	52.5
168	183	57.1	62.4	239	75.8	149	46.7	225	339	94.7	163	158	134	225
0.22	0.12	b.d.	0.24	0.18	0.2	0.36	b.d.	b.d.	b.d.	b.d.	0.3	0.34	0.39	0.35
0.62	1.14	0.41	0.62	0.42	1.9	1.58	0.17	0.07	0.25	0.21	0.18	0.2	0.22	0.2
0.34	0.27	0.23	0.17	0.11	0.2	0.34	0.13	0.06	0.09	0.07	0.06	0.08	0.07	0.07
315	376	283	248	267	111	414	306	199	253	253	304	314	315	346
15.99	11.44	22.96	17.12	16.33	34.7	15.98	31.07	16.31	28.52	27.98	34.41	34.92	34.81	40.44
31.5	33.2	46.86	40.7	32.1	82.5	75.4	79.77	40.93	76.38	73.33	83.6	92.3	84.4	104.2
4.12	5.78	2.49	3.72	3.31	9.90	5.54	2.59	1.17	3.72	2.66	2.71	3.05	3.10	3.24
8.99	12.22	7.05	8.94	7.74	24.80	13.26	8.16	3.83	10.45	8.35	8.56	9.57	9.52	10.10
1.25	1.62	1.24	1.32	1.11	3.37	1.84	1.58	0.73	1.73	1.50	1.61	1.70	1.68	1.88
6.13	6.86	7.00	6.30	5.66	17.40	8.15	8.68	4.25	8.97	8.17	9.04	9.51	9.61	10.85
1.78	1.75	2.43	1.93	1.85	4.50	2.27	3.07	1.60	3.09	2.88	3.30	3.31	3.40	3.96
0.61	0.58	0.94	0.70	0.66	1.59	0.94	1.21	0.68	1.55	1.11	1.19	1.25	1.22	1.37
2.06	1.88	3.56	2.64	2.45	5.68	2.77	4.63	2.40	4.13	4.09	4.91	4.87	4.80	5.73
0.38	0.31	0.62	0.48	0.46	0.89	0.47	0.83	0.42	0.78	0.76	0.90	0.89	0.92	1.01
2.50	1.88	4.26	2.89	2.79	5.50	2.99	5.36	2.80	4.93	4.72	5.84	5.85	5.82	6.80
0.57	0.41	0.92	0.64	0.62	1.15	0.63	1.20	0.63	1.12	1.10	1.29	1.27	1.28	1.51
1.76	1.26	2.59	1.91	1.80	3.47	1.83	3.41	1.84	3.02	3.05	3.82	3.84	3.70	4.52
0.25	0.19	0.35	0.26	0.29	0.47	0.28	0.53	0.27	0.45	0.45	0.57	0.59	0.57	0.68
1.57	1.27	2.07	1.78	1.76	3.33	1.81	3.34	1.69	2.72	2.77	3.56	3.66	3.53	4.09
0.24	0.19	0.31	0.29	0.27	0.54	0.28	0.52	0.24	0.43	0.44	0.55	0.59	0.53	0.67
52.2	46.8	57.4	61.0	69.1	59.4	50.6	56.3	65.7	58.5	59.7	60.1	53.6	52.0	52.9
0.3	0.2	0.3	0.3	0.4	0.3	0.2	0.7	0.8	0.7	0.6	0.7	0.7	0.7	0.8
5.8	4.2	2.5	3.0	4.4	1.9	1.4	1.2	1.0	0.9	0.8	1.1	1.2	1.0	1.1
0.63	0.67	0.69	0.75	0.62	0.65	1.18	0.92	0.91	0.88	0.90	0.90	0.99	0.88	0.94
3.2	5.6	1.5	2.5	2.3	3.6	3.7	0.9	0.8	1.7	1.2	0.9	1.0	1.1	1.0

Table DR1 (cont'd)

VL01A 078	VL01A 301b	VL01A 331b	VL02A 021	VL02A 028	VL02A 080	VL02A 024	VL02A 274	RAX01 099	RAX01 047	VL02A 087	VL01A 133b	VL02A 197	VL02A 084	RAX01 048
445059	465984	467424	443604	464567	467364	447910	484804	501089	503114	467315	446907	480124	468133	503137
5338271	5358028	5360730	5337182	5354343	5362844	5341643	5377692	5383217	5387085	5362451	5340466	5372628	5363329	5387255
12A/04	12A/06	12A/06	12A/04	12A/06	12A/06	12A/04	12A/11	12A/10	12A/10	12A/06	12A/04	12A/11	12A/06	12A/10
HR <sub>1</sub>	HR <sub>2</sub>	HR <sub>2</sub>	HR <sub>2</sub>	HR <sub>3</sub>	HR <sub>3</sub>	HR <sub>3</sub>	HR <sub>4</sub>							
diab	bslt	gabb	bslt	pbslt	bslt	pbslt	bslt	mtuff	pbslt	bslt	mtuff	bslt	bslt	pbslt
5	5	6	3	3	3	3	3	2	2	3	5	3	6	2
47.99	47.94	49.21	47.61	47.73	50.17	50.22	47.71	51.17	50.00	49.25	48.17	46.78	48.29	50.14
1.81	1.73	1.77	1.46	1.39	1.45	1.43	1.97	1.26	1.90	1.65	0.73	1.24	1.26	0.55
13.04	14.63	13.43	14.76	14.79	13.80	14.21	13.72	15.48	14.51	14.61	16.36	16.42	14.46	15.35
13.62	13.32	13.01	10.54	12.81	12.00	11.71	13.92	9.72	12.99	11.56	9.96	11.41	11.71	8.61
0.24	0.21	0.19	0.19	0.21	0.21	0.20	0.24	0.16	0.21	0.21	0.18	0.18	0.19	0.14
7.71	7.48	6.75	7.48	8.16	7.58	6.18	7.23	6.64	5.55	7.16	8.33	7.20	7.43	9.42
8.51	8.67	10.43	13.71	9.15	10.26	9.60	10.32	6.64	7.76	11.23	11.24	11.56	11.44	7.57
3.69	3.91	3.22	2.13	3.43	3.00	4.67	2.84	5.99	4.74	2.76	2.98	2.78	2.61	4.89
0.60	0.04	0.10	0.04	0.20	0.49	0.05	0.06	0.10	0.24	0.07	0.31	0.14	0.10	0.12
0.14	0.12	0.15	0.14	0.11	0.13	0.13	0.18	0.14	0.18	0.18	0.04	0.09	0.09	0.09
2.60	2.68	1.50	2.18	2.90	1.14	2.12	2.26	3.56	3.29	2.04	2.66	2.65	2.20	4.34
99.95	100.73	99.80	100.31	100.95	100.30	100.60	100.52	100.86	101.36	100.77	100.96	100.53	99.81	101.23
223	13.6	39.0	28.1	70.8	27.5	32.6	17.4	23.7	97.6	15.3	48.7	29.9	20.0	76.8
103	154	151	151	231	199	196	163	227	194	143	337	330	116	327
0.46	0.16	0.20	0.34	0.45	2.5	0.60	0.19	0.08	0.38	0.67	0.44	0.30	0.40	0.15
3.20	2.88	2.90	2.70	2.30	2.40	2.40	3.50	2.15	3.52	2.80	0.94	1.90	2.10	0.88
4.03	2.90	3.00	5.10	2.03	3.20	2.31	3.94	4.11	3.53	7.93	0.29	1.12	2.60	0.86
50	72	27	69	107	84	77	70	83	102	65	131	115	35	143
0.83	1.57	1.2	0.5	0.9	4.6	0.6	1.9	2.04	1.6	2.5	0.6	3.7	0.2	1.14
8.19	0.56	0.9	0.42	2.68	29.4	0.46	0.36	1.37	3.5	0.74	7.58	2.46	2.1	1.41
44.3	50.3	40	38	43	40	45	46.6	43	34.1	47.0	37.3	44.2	39	35.7
134	143	102	300	91.4	125	167	117	159	140	277	83.8	312	162	115
b.d.	b.d.	0.2	0.48	0.31	0.34	0.34	0.38	b.d.	b.d.	0.67	b.d.	0.18	b.d.	b.d.
0.3	0.21	0.3	0.36	0.17	0.23	0.18	0.34	0.65	0.53	0.46	b.d.	0.11	b.d.	0.6
0.1	0.12	0.1	0.11	0.05	0.08	0.07	0.10	0.54	0.16	0.18	0.01	0.03	b.d.	0.23
351	351	354	235	281	303	314	406	316	514	298	225	286	307	212
40.74	41.63	42.5	33.41	35.51	33.66	35.14	46.87	26.35	44.58	34.87	19.79	29.85	30.7	12.22
118.87	100.13	94	106	78.8	91.1	89.7	126.3	72.66	113.98	115.7	23.27	74.3	61	23.58
4.51	3.43	4.00	5.00	2.54	3.51	3.16	4.61	5.15	5.54	6.58	0.50	1.90	2.80	2.52
13.39	10.67	11.50	14.09	8.06	10.56	9.50	14.41	12.36	15.41	17.18	1.89	7.01	8.70	6.18
2.36	1.93	2.09	2.26	1.48	1.75	1.71	2.47	1.89	2.58	2.54	0.44	1.38	1.40	0.95
12.44	10.57	11.80	12.10	8.29	9.68	9.47	13.12	9.28	13.28	12.58	2.95	7.76	8.30	4.68
4.19	3.99	4.10	3.99	3.19	3.37	3.43	4.80	2.81	4.54	4.08	1.48	3.17	2.40	1.54
1.41	1.62	1.32	1.46	1.15	1.21	1.30	1.54	1.05	1.78	1.47	0.66	1.18	1.14	0.59
5.74	5.76	5.90	5.38	4.87	4.67	4.97	6.40	3.92	6.40	5.12	2.41	4.27	4.95	1.91
1.04	1.09	1.05	0.90	0.89	0.85	0.93	1.20	0.70	1.16	0.88	0.48	0.76	0.79	0.36
6.80	7.00	6.50	5.83	5.74	5.82	5.76	7.89	4.61	7.83	5.76	3.40	5.00	5.24	2.27
1.50	1.62	1.40	1.25	1.29	1.22	1.32	1.71	1.00	1.72	1.21	0.78	1.10	1.06	0.49
4.29	4.55	4.17	3.76	3.74	3.53	3.93	5.27	2.89	5.00	3.56	2.22	3.35	3.22	1.42
0.63	0.69	0.64	0.53	0.55	0.48	0.54	0.75	0.45	0.76	0.52	0.34	0.50	0.49	0.22
4.04	4.22	4.01	3.20	3.76	3.60	3.62	4.74	2.80	4.76	3.17	2.11	2.93	2.90	1.33
0.65	0.68	0.68	0.50	0.53	0.53	0.59	0.73	0.43	0.76	0.51	0.34	0.46	0.45	0.23
55.2	55.0	53.1	60.7	58.1	57.9	53.5	53.1	59.8	48.2	57.4	64.6	57.9	58.0	70.5
0.7	0.8	0.6	0.7	0.7	0.7	0.8	0.7	0.4	0.5	0.7	n.a.	0.8	n.a.	0.2
1.0	1.1	1.2	0.9	1.2	1.0	1.3	1.1	1.2	1.5	0.8	1.6	1.6	1.0	2.7
1.01	0.89	0.81	0.94	0.88	0.96	0.93	0.94	0.92	0.89	1.01	0.56	0.83	0.90	0.54
1.4	1.0	1.2	1.9	0.8	1.2	1.1	1.2	2.2	1.4	2.5	0.3	0.8	1.2	2.3

Table DR1 (cont'd)

RAX01 032	RAX01 033	VL01A 080a	RAX00 709	RAX00 706	RAX00 707	RAX01 054	RAX01 062	RAX01 061	RAX01 060	RAX01 067	RAX01 076	RAX01 064	RAX01 035	RAX01 056
496519	497088	446204	443569	440819	440869	488958	534506	533826	532918	529249	508414	530989	510647	502988
5379867	5379494	5340308	5336518	5336018	5336018	5379606	5407801	5407267	5406265	5401639	5394308	5406069	5395316	5387700
12A/11	12A/11	12A/04	12A/04	12A/04	12A/04	12A/11	12A/15	12A/15	12A/15	12A/15	12A/10	12A/15	12A/10	12A/10
HR <sub>4</sub>	HR <sub>4</sub>	HR <sub>4</sub>	HR <sub>5</sub>											
mtuff	pbslt	bslt	bslt	pbslt	pbslt	and	pbslt	pbslt	pand	pbslt	mtuff	pbslt	bslt	mtuff
2	2	5	1	1	1	2	2	2	2	2	2	2	2	2
40.89	44.35	52.66	57.40	47.20	44.40	56.89	49.97	41.80	58.61	48.46	50.37	49.73	48.74	51.25
0.62	0.82	0.63	0.64	0.48	0.54	0.73	1.51	0.80	0.62	0.74	0.95	0.71	0.91	0.75
14.27	17.81	16.14	17.30	16.00	18.20	17.20	16.24	14.30	11.56	14.56	14.85	13.80	15.47	16.28
8.68	10.16	7.95	8.07	6.68	9.10	8.62	8.93	7.86	10.06	8.92	8.09	7.04	8.76	8.14
0.15	0.25	0.10	0.13	0.18	0.23	0.17	0.14	0.13	0.50	0.15	0.17	0.18	0.15	0.23
5.09	6.84	9.93	4.39	11.00	9.80	3.25	5.97	5.87	7.95	7.80	4.33	5.28	8.94	8.29
12.52	6.66	3.07	2.59	7.08	8.16	6.60	5.24	12.55	4.70	10.00	8.22	10.28	5.84	4.84
1.70	3.92	4.82	3.57	2.43	1.91	2.44	5.27	3.74	0.12	3.82	4.13	4.37	3.61	4.83
3.11	2.01	0.04	1.34	0.07	0.39	1.69	0.46	0.16	0.19	1.37	0.46	0.22	0.42	1.73
0.05	0.09	0.09	0.09	0.24	0.27	0.24	0.26	0.18	0.09	0.27	0.24	0.14	0.20	0.16
12.96	8.55	5.53	7.21	12.64	11.91	3.23	7.26	13.72	5.45	5.30	9.40	9.19	8.30	4.74
100.06	101.46	100.96	99.00	98.20	99.30	101.07	101.25	101.11	99.85	101.39	101.20	100.93	101.35	101.25
234	152	77.4	337	95.0	102	251	348	67.1	706	504	140	229	92.7	359
501	75	64	19	1160	1040	5	81	323	35	303	55	208	289	59
0.99	1.4	0.37	1.3	0.12	0.36	1.6	0.96	0.77	0.96	0.38	2.0	0.11	0.83	1.0
1.24	1.33	0.96	2.50	1.10	1.20	2.77	3.97	2.32	1.01	2.14	2.89	1.34	1.91	1.91
0.60	0.76	0.92	2.40	1.80	2.00	2.47	7.38	3.43	1.29	4.35	6.07	1.50	3.87	3.40
195	39	60	b.d.	388	341	b.d.	8	156	40	81	36	92	132	50
1.18	1.06	5.43	2	7	13	4.4	2.7	0.99	6.68	2.1	5.14	0.95	2.42	1.73
45.1	44.1	0.53	26.0	1.2	9.1	59.9	9.33	3.2	5.45	32.3	14.7	4.05	11.9	33.0
32.5	39.7	34	22	31	36	17.7	19.7	31.7	24.9	37.6	26.1	36	31.3	36.2
229	93.3	166	112	343	498	451	205	169	121	165	153	157	188	173
b.d.	b.d.	b.d.	0.2	0.16	0.15	b.d.	0.4	b.d.						
1.25	1.2	0.72	4.2	1.9	2.1	10.39	4.42	3.6	1.82	5.08	5.62	1.64	3.26	3.75
0.39	0.44	0.29	1.3	0.74	0.68	3.51	1.4	1.19	0.49	2.56	0.98	0.54	0.58	0.75
223	343	278	153	254	434	168	179	223	231	242	244	257	266	316
15.46	16.81	14.83	22	16	18	17.31	22.3	16.29	12.08	16.18	20.4	12.07	15.05	15.58
37.42	43.96	30.28	88	34	40	94.53	153.56	87.57	34.36	74.37	103.18	43.88	66.61	66.25
4.75	4.09	3.70	11.00	8.50	9.90	26.98	19.28	13.00	7.27	11.79	14.42	7.34	10.92	12.12
11.23	10.32	9.16	23.00	18.00	21.00	56.65	42.97	28.60	15.16	26.24	31.99	15.60	25.08	25.84
1.72	1.53	1.38	2.90	2.50	2.80	7.16	5.59	3.78	1.99	3.66	4.35	2.15	3.38	3.48
8.09	6.91	6.34	12.00	11.00	12.00	28.48	22.46	15.57	8.64	16.45	18.88	9.20	13.78	14.51
2.26	2.04	1.94	2.60	2.60	2.90	5.53	4.52	3.39	1.96	3.54	3.93	2.31	3.32	3.08
0.85	0.81	0.69	0.94	0.83	0.90	1.68	1.46	0.98	0.73	1.00	1.22	0.78	1.06	1.07
2.52	2.62	2.21	2.90	2.70	3.10	4.47	4.40	3.09	2.30	3.16	3.79	2.30	2.96	3.07
0.47	0.45	0.38	0.54	0.46	0.51	0.63	0.68	0.50	0.36	0.48	0.59	0.37	0.50	0.48
2.88	2.91	2.58	3.20	2.70	3.00	3.46	4.05	2.92	2.27	3.02	3.72	2.26	2.97	3.01
0.62	0.64	0.56	0.70	0.55	0.60	0.72	0.86	0.61	0.50	0.60	0.81	0.48	0.62	0.64
1.75	1.82	1.70	1.90	1.40	1.60	1.99	2.40	1.91	1.44	1.71	2.33	1.38	1.78	1.82
0.26	0.27	0.25	0.33	0.23	0.26	0.30	0.37	0.27	0.21	0.25	0.35	0.21	0.27	0.27
1.64	1.74	1.64	2.30	1.60	1.70	1.98	2.22	1.78	1.29	1.49	2.27	1.38	1.69	1.69
0.26	0.26	0.26	0.39	0.23	0.28	0.32	0.35	0.29	0.21	0.24	0.36	0.21	0.27	0.27
56.1	59.5	73.1	54.2	78.2	70.1	45.1	59.3	61.9	63.3	65.6	53.8	62.0	69.0	68.9
0.2	0.2	0.2	0.1	0.2	0.2	0.1	0.2	0.2	0.2	0.1	0.1	0.2	0.2	0.2
7.4	5.0	3.7	4.3	4.4	4.6	10.2	2.4	3.5	5.3	2.5	2.2	4.6	2.6	3.3
0.59	0.77	0.55	1.20	0.46	0.49	0.61	1.21	0.92	0.62	0.75	0.93	0.68	0.71	0.76
3.5	2.9	2.8	5.8	6.5	7.1	16.6	10.6	8.9	6.9	9.7	7.7	6.5	7.9	8.7

Table DR1 (cont'd)

RAX01 049	RAX01 031	RAX01 063	RAX01 058	RAX01 065	RAX02 129	RAX02 103	RAX02 153	RAX02 154	RAX02 102	RAX02 101	VL02A 014	RAX00 710	RAX01 057	RAX01 910
501727	500049	535078	503298	524147	537110	514047	513134	509807	513531	513126	440325	443569	503226	502989
5386376	5384112	5409320	5388722	5404056	5410490	5397396	5403769	5402136	5396921	5396703	5335838	5336518	5388500	5387699
12A/10	12A/10	12A/15	12A/10	12A/15	12A/16	12A/10	12A/15	12A/15	12A/10	12A/10	12A/04	12A/04	12A/10	12A/10
HR <sub>5</sub>	HBF	HBF	HBF	HBF										
mtuff	bslt	pbslt	diab	gabb	pbslt	bslt	bslt	pbslt	bslt	bslt	ituff	ftuff	ftuff	ftuff
2	2	2	2	2	4	4	4	4	4	6	6	1	2	2
51.10	49.36	48.26	51.83	49.64	52.48	49.98	52.73	48.13	50.65	55.84	48.21	65.60	77.21	76.04
0.89	0.78	0.79	0.83	1.69	0.69	1.08	0.72	0.87	1.16	1.21	1.03	0.43	0.14	0.18
16.51	16.69	16.54	14.92	15.71	14.10	19.12	16.64	18.69	16.49	14.86	16.41	15.10	12.48	12.60
9.46	8.33	10.06	11.90	13.46	7.68	12.18	9.61	9.88	12.18	11.22	7.45	6.34	1.45	1.27
0.24	0.08	0.16	0.17	0.23	0.14	0.13	0.18	0.24	0.20	0.12	0.12	0.14	0.05	0.18
7.29	9.58	5.94	7.34	5.42	5.04	4.02	3.73	9.10	6.87	3.92	5.64	2.91	1.55	1.69
6.41	5.18	8.48	2.58	6.11	10.42	2.41	9.28	3.18	4.13	5.63	8.83	1.74	0.20	0.32
4.96	1.44	5.28	3.96	4.67	2.73	5.78	1.77	4.66	4.15	3.29	4.88	4.96	1.16	1.17
0.46	5.34	0.09	1.57	1.32	1.28	2.31	2.32	0.71	0.80	0.44	0.45	0.45	4.70	4.74
0.16	0.13	0.17	0.20	0.22	0.22	0.26	0.15	0.16	0.27	0.29	0.36	0.06	0.03	0.04
3.76	4.53	5.50	5.80	2.91	5.12	3.56	3.55	5.14	3.92	2.90	6.40	4.51	2.30	2.37
101.25	101.46	101.27	101.10	101.40	99.98	100.86	100.72	100.81	100.86	99.75	99.85	99.80	101.28	100.62
129	464	79.1	253	462	726	409	507	497	300	169	233	162	241	635
109	223	38	21	5	420	54	53	152	21	27	157	15	b.d.	b.d.
0.62	2.5	0.06	1.0	0.14	0.41	2.5	0.81	0.50	0.27	1.0	0.50	0.40	2.0	1.7
1.80	1.80	1.76	1.62	2.61	2.30	2.40	1.90	2.10	2.50	2.70	2.70	2.40	3.70	4.18
2.87	3.25	3.08	1.81	2.74	4.53	4.13	2.43	2.87	4.50	4.30	11.30	2.10	9.59	9.36
55	88	36	22	7	132	18	24	59	19	10	104	b.d.	b.d.	b.d.
3.09	1.26	5.03	1.82	4.01	16	5.4	9.5	4.3	7.6	2.9	1.3	8	1.18	13.71
7.24	124	1.59	19.4	31.2	37.0	60.8	55.6	11.6	9.7	19.2	11.5	6.4	85.5	69.3
33.3	40.8	31.1	17.1	38.1	28.0	40.2	38.9	52.5	42.3	31	27	15	6.68	b.d.
236	191	122	137	275	592	76.7	312	396	154	382	184	184	2.5	8.22
b.d.	b.d.	b.d.	b.d.	b.d.	0.33	0.38	0.22	0.37	0.42	0.3	0.5	0.2	0.6	0.66
3.57	2.86	3.52	1.64	3.69	6.12	4.66	4.77	3.44	4.9	5.3	10.8	4.2	8.44	8.38
0.69	0.54	1.02	0.72	0.78	0.91	0.37	1.03	0.84	0.88	1.1	2.1	1.2	2.06	2.17
319	320	463	553	751	218	229	315	315	357	383	256	90	9	9
16.4	15.97	16.48	16.29	25.97	20.14	26.04	16.82	17.04	24.21	28.3	21.4	17	27.91	12.26
58.51	61.6	61.2	50.22	92.61	89.8	85.5	72.5	77.2	91.2	87.1	111.8	87	107.92	133.3
13.98	10.87	13.69	6.52	15.73	19.34	18.48	16.89	13.05	20.14	23.00	18.10	10.00	23.86	17.06
28.98	22.75	27.02	14.87	34.32	38.03	38.22	34.19	27.64	43.44	49.20	41.30	21.00	50.17	34.28
3.62	3.02	3.43	2.06	4.72	4.79	5.44	4.35	3.59	6.00	6.25	5.49	2.50	6.16	4.05
15.21	12.62	14.39	9.39	20.10	19.08	22.65	16.95	14.39	25.24	27.30	24.20	9.90	22.97	14.17
3.48	2.87	3.24	2.37	4.93	4.02	5.33	3.60	3.18	5.50	5.70	4.30	2.20	4.66	2.60
1.18	0.98	1.08	0.73	1.72	1.08	1.59	1.08	1.01	1.37	1.83	1.33	0.69	0.72	0.42
3.53	2.86	3.41	2.65	4.91	3.89	5.17	3.53	3.34	5.30	5.61	4.37	2.50	4.42	2.30
0.55	0.48	0.50	0.43	0.79	0.56	0.81	0.51	0.53	0.81	0.74	0.55	0.43	0.72	0.36
3.30	2.86	2.98	2.87	4.84	3.49	4.76	2.86	3.12	4.66	5.12	3.60	2.60	4.69	2.10
0.68	0.63	0.64	0.62	1.01	0.72	1.08	0.60	0.61	1.01	1.03	0.72	0.57	1.03	0.45
1.83	1.62	1.86	1.83	2.97	2.06	2.98	1.73	1.89	2.84	2.83	2.09	1.60	2.96	1.35
0.28	0.27	0.27	0.27	0.42	0.31	0.47	0.25	0.26	0.44	0.44	0.27	0.27	0.46	0.22
1.71	1.61	1.89	1.68	2.70	2.05	2.73	1.56	1.66	2.66	2.95	2.21	2.00	2.77	1.55
0.27	0.26	0.27	0.25	0.40	0.32	0.43	0.25	0.27	0.43	0.35	0.36	0.33	0.43	0.24
62.7	71.5	56.3	57.3	46.7	58.9	41.8	45.8	66.8	55.1	43.2	62.3	50.0	70.0	74.4
0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.1	0.1	0.1	0.1
4.5	3.1	4.1	3.4	5.4	4.0	4.2	6.5	4.2	4.2	5.0	1.5	4.4	2.3	1.7
0.60	0.76	0.67	0.75	0.67	0.79	0.57	0.72	0.86	0.59	0.54	0.92	1.41	0.82	1.82
10.0	8.2	8.8	4.7	7.1	11.5	8.3	13.2	9.6	9.2	9.5	10.0	6.1	10.5	13.4

Table DR1 (cont'd)

RAX01 051	RAX01 011	RAX01 030	RAX01 055	RAX01 908	RAX01 009	RAX01 081	RAX01 909	RAX02 155	RAX02 144	RAX02 143	RAX02 152	VL01A 308	VL01A 229	VL01A 292b
500800	506205	503767	503672	506248	506539	505297	503520	509807	529723	529793	513700	466576	446955	465233
5385042	5388840	5386729	5387478	5392154	5391273	5387579	5389050	5402136	5401170	5401149	5404021	5358915	5340646	5359136
12A/10	12A/15	12A/15	12A/15	12A/15	12A/06	12A/04	12A/06							
HBF														
ftuff	fdyke	rhyo	gran	ftuff	ftuff	ftuff	ftuff							
2	2	2	2	2	2	2	2	6	4	4	4	5	5	5
79.38	74.23	67.48	68.93	69.28	65.21	63.96	63.75	81.42	77.60	69.36	63.52	73.89	73.53	75.56
0.26	0.30	0.44	0.45	0.50	0.49	0.50	0.58	0.11	0.24	0.39	0.81	0.18	0.30	0.28
11.33	13.65	15.88	15.87	14.52	17.11	18.45	16.25	9.87	10.82	14.78	16.23	13.60	12.96	11.93
2.15	2.60	4.54	4.42	5.03	5.68	5.74	7.00	1.52	2.58	3.95	4.95	2.11	2.42	2.23
0.46	0.06	0.08	0.07	0.09	0.09	0.10	0.10	0.03	0.07	0.08	0.08	0.14	0.08	0.07
0.33	1.35	1.85	1.82	2.04	2.62	2.41	3.10	0.65	1.41	1.83	2.71	2.93	0.63	1.57
0.03	1.49	0.37	0.56	1.35	1.02	0.30	1.37	0.78	1.53	1.99	1.46	0.26	1.40	1.12
2.76	2.77	3.89	3.22	2.47	3.56	3.08	3.28	4.09	4.70	4.65	6.78	0.82	4.04	2.07
2.59	2.56	2.77	2.97	2.60	2.45	3.47	2.61	0.62	0.29	1.30	0.71	3.83	2.63	2.30
0.04	0.04	0.08	0.10	0.12	0.12	0.19	0.11	0.02	0.05	0.08	0.23	0.03	0.05	0.07
1.58	1.85	2.56	2.47	2.79	2.87	3.14	2.95	0.70	0.85	1.80	2.59	2.71	1.83	3.06
100.92	100.91	99.94	100.88	100.79	101.21	101.36	101.09	99.87	100.16	100.23	100.09	100.49	99.87	100.25
697	796	591	588	710	533	684	644	540	339	280	268	1036	806	643
2	13	17	17	42	33	26	57	27	46	37	20	2	1	2
1.8	2.9	2.6	3.0	2.4	2.9	3.2	2.0	b.d.	0.12	0.84	0.74	1.8	1.6	1.4
3.36	3.42	4.02	4.03	3.88	4.01	3.84	3.72	2.20	3.30	3.00	4.40	3.57	3.07	3.50
6.70	12.82	8.13	8.23	6.59	8.85	8.73	7.42	6.10	5.25	5.70	6.06	7.77	9.73	6.14
b.d.	7	9	9	15	14	13	25	5	8	8	6	b.d.	b.d.	b.d.
9.19	4.13	7.15	4.74	13.49	27.59	114.03	11.94	7.4	3.8	8	20.6	4.2	31.54	3.04
82.2	71.7	88.9	92.4	78.3	81.9	113	76.2	14.0	4.57	37.4	20.0	91.0	70.9	66.2
5.95	7.77	15.05	14.5	13.8	18.5	18.9	20.6	4	10.4	13.5	21.9	3.63	9.55	9.56
58.9	257	42.8	88.7	164	148	47.6	156	172	133	207	143	32.1	186	136
0.42	1.04	0.46	0.48	0.36	0.61	0.53	0.39	0.6	0.48	0.48	0.43	0.58	0.58	0.38
10.12	22.59	10.29	10.85	9.24	11.24	10.77	9.63	8.1	6.21	4.49	10.81	11.73	10.05	7.15
2.57	5.36	2.32	2.57	2.17	2.67	2.63	2.06	1.4	1.56	1.16	2.39	1.58	3.14	1.76
32	35	75	78	90	93	94	108	8	46	65	66	24	28	30
19.16	15.27	22.6	20.53	21.95	23.3	26.44	22.86	14.9	19.85	16.01	35.74	14.11	26.39	19.03
110.56	108.93	137.93	138.27	146.45	132.46	120.32	131.58	65.9	123	122.7	178	114.96	101.02	127.98
26.57	32.63	28.27	22.99	25.40	23.38	23.41	26.26	18.80	16.16	18.51	42.88	18.75	27.11	20.09
46.44	66.63	43.28	40.16	49.16	55.54	49.05	53.98	36.30	32.48	30.05	85.37	38.54	52.33	40.92
5.63	6.18	6.07	5.44	6.08	5.58	6.07	6.36	3.81	3.62	3.85	11.17	3.89	6.45	4.92
20.10	18.73	21.91	20.52	22.60	20.08	22.39	24.08	14.40	13.63	14.12	43.26	12.25	22.64	18.01
3.67	3.17	4.08	4.17	4.39	4.15	4.79	4.76	2.50	2.79	2.75	8.14	2.13	4.58	3.67
0.85	0.66	1.02	1.04	1.23	1.02	1.07	1.25	0.38	0.61	0.82	2.07	0.40	0.98	0.87
3.36	2.38	3.82	3.88	4.12	3.90	4.52	4.31	1.88	2.74	2.63	7.52	1.77	4.24	3.31
0.54	0.41	0.63	0.64	0.63	0.71	0.77	0.67	0.28	0.48	0.41	1.05	0.32	0.72	0.55
3.20	2.54	3.94	3.75	3.80	4.06	4.71	4.07	2.50	3.00	2.46	5.75	2.10	4.36	3.27
0.68	0.55	0.84	0.83	0.81	0.88	1.04	0.87	0.49	0.71	0.52	1.25	0.49	0.95	0.72
1.98	1.76	2.48	2.56	2.28	2.62	3.07	2.53	1.63	2.14	1.49	3.58	1.53	2.70	2.08
0.32	0.30	0.40	0.41	0.34	0.42	0.47	0.38	0.21	0.35	0.23	0.55	0.28	0.41	0.33
2.17	2.14	2.55	2.73	2.34	2.76	3.16	2.56	2.11	2.20	1.55	3.74	1.90	2.67	2.08
0.36	0.33	0.41	0.45	0.37	0.42	0.49	0.38	0.30	0.38	0.25	0.56	0.34	0.44	0.35
31.8	53.1	47.0	47.3	46.9	50.1	47.8	49.1	48.2	54.4	50.2	54.4	75.2	36.2	60.5
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.2	0.2	0.1	0.1	0.1
3.7	2.4	3.2	2.6	3.6	2.5	2.5	3.3	2.9	2.9	3.0	6.6	2.2	2.6	3.0
1.07	1.22	1.20	1.18	1.19	1.13	0.89	0.98	0.94	1.57	1.59	0.78	1.92	0.78	1.24
14.9	18.6	13.5	10.3	13.2	10.3	9.0	12.5	10.9	9.0	14.6	14.0	12.0	12.4	11.8

Table DR1 (cont'd)

VL01A 309	VL01A 360	VL01A 133a	VL01A 313	VL02A 173	VL02A 283	VL02A 013b	VL02A 158	VL02A 022b	VL02A 295a	VL02A 012	VL02A 081
466970	440098	446907	467207	474579	481058	440098	470197	446730	484785	439962	468746
5359300	533591	5340466	5359806	5368625	5373193	533591	5365831	5340544	5377776	5335514	5364078
12A/06	12A/04	12A/04	12A/06	12A/06	12A/11	12A/04	12A/06	12A/04	12A/11	12A/04	12A/06
HBF	HBF	HBF	HBF	HBF	HBF	HBF	HBF	HBF	HBF	HBF	HBF
ftuff	ftuff	ftuff	ftuff	gran	ftuff	ftuff	ftuff	ftuff	ftuff	ftuff	ftuff
6	5	3	3	3	3	6	3	6	3	3	3
74.80	70.43	74.16	75.05	75.43	76.19	75.13	70.56	74.28	70.28	77.43	64.45
0.33	0.34	0.26	0.33	0.28	0.35	0.33	0.34	0.37	0.39	0.17	0.37
12.24	14.78	13.26	11.87	12.79	8.28	11.81	15.53	12.05	14.99	12.78	15.55
3.34	2.97	2.44	3.22	2.09	7.06	3.48	2.98	3.82	3.79	1.02	5.80
0.05	0.07	0.19	0.07	0.12	0.60	0.06	0.05	0.06	0.08	0.05	0.10
1.15	1.74	0.96	1.17	0.54	1.16	1.12	1.07	1.10	1.41	0.56	2.60
0.86	0.31	1.14	1.45	1.50	0.82	0.59	0.67	0.70	0.58	0.34	2.76
3.63	2.58	4.68	2.48	4.81	1.95	2.58	3.08	2.53	3.21	4.21	5.82
1.55	4.09	1.84	2.57	1.72	2.58	2.53	3.80	2.97	3.11	2.26	0.48
0.05	0.06	0.06	0.06	0.04	0.07	0.05	0.07	0.04	0.08	0.03	0.07
1.80	2.54	1.09	1.82	0.57	0.68	2.10	1.66	1.90	1.89	1.18	2.09
99.86	99.92	100.09	100.11	99.90	99.76	99.86	99.82	99.90	99.81	100.04	100.10
470	877	396	912	821	305	647	846	619	763	521	206
3	8	47	46	35	72	7	32	7	34	29	26
0.90	3.1	0.90	2.1	2.4	11.9	1.7	3.9	2.1	3.4	1.4	0.67
3.10	4.24	3.50	3.40	3.40	1.70	4.10	3.70	3.00	3.90	4.40	2.00
6.70	9.16	7.57	6.80	7.54	6.82	7.30	9.39	7.40	8.88	18.52	4.74
3	b.d.	8	7	5	31	6	8	5	11	6	11
4.4	17.75	23.7	29	20	5.4	8.5	42	4.5	19.9	4.7	11.1
46.1	123	52.5	72.0	63.8	167	75.5	136	89.6	117	81.6	10.5
8	12.3	16.3	15.1	9.97	11.8	8	15.2	9	18.1	5	23.1
89.2	48.8	93.6	145	168	101	60.3	104	50.7	146	132	283
0.4	0.57	0.7	0.61	0.46	0.56	0.4	0.6	0.5	0.77	1.27	0.52
9.1	9.73	8.05	7.96	9.26	5.25	7.7	11.46	7.7	9.28	14.11	4.79
1.9	2.54	1.86	1.93	1.88	2.47	1.9	2.84	2.5	2.21	3.24	1.24
44	45	46	73	25	47	49	56	61	71	109	139
21.9	25.45	25.46	21.41	22.69	21.46	16.2	29.19	22.9	27.98	27.6	16.12
111	154.16	129.5	131.8	119.4	60.5	134.6	126.9	135.6	126	113.5	80.8
28.20	26.52	24.15	24.66	29.80	23.54	26.10	34.49	26.90	27.83	37.08	16.22
48.50	51.32	46.80	46.90	58.16	54.39	48.10	65.29	50.20	52.75	78.29	32.20
5.59	6.33	5.56	5.70	6.85	5.49	5.38	7.63	5.87	6.44	9.50	3.61
21.40	22.08	20.43	20.63	25.27	21.55	20.80	26.93	21.40	23.61	35.39	13.69
3.90	4.48	4.03	3.88	4.60	4.63	3.60	5.23	3.50	4.87	6.61	2.96
0.89	1.03	0.77	0.93	1.19	1.01	0.72	1.04	0.86	1.03	1.15	0.77
3.31	4.09	4.02	3.59	3.98	5.01	2.62	4.69	3.51	4.76	5.17	2.61
0.51	0.67	0.64	0.57	0.61	0.76	0.49	0.78	0.49	0.73	0.71	0.41
3.70	4.19	4.12	3.48	3.66	4.46	3.00	4.61	4.18	4.43	4.44	2.54
0.67	0.92	0.85	0.74	0.74	0.78	0.62	1.05	0.69	1.01	1.00	0.54
2.15	2.61	2.65	2.25	2.28	2.26	1.54	3.07	2.30	2.88	3.21	1.71
0.38	0.42	0.44	0.35	0.35	0.36	0.30	0.42	0.33	0.44	0.53	0.26
2.04	2.70	2.72	2.27	2.37	2.06	2.23	2.94	2.89	2.95	3.65	1.70
0.30	0.46	0.44	0.38	0.39	0.33	0.42	0.49	0.36	0.44	0.58	0.28
42.9	56.1	46.2	44.2	36.0	26.4	41.2	43.9	38.6	44.8	54.5	49.4
0.1	0.1	0.1	0.1	0.2	0.2	0.2	0.1	0.2	0.1	0.1	0.2
3.9	2.7	3.0	3.4	3.7	3.2	3.3	3.4	3.4	2.9	1.9	3.2
1.01	1.22	1.14	1.21	0.92	0.46	1.33	0.86	1.38	0.92	0.61	0.97
16.9	12.0	10.8	13.3	15.3	13.9	14.3	14.3	11.4	11.5	12.4	11.6

Table DR1 (cont'd)

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