

Expedition 320 Site U1334, Table T22. Magnetostratigraphy, Site U1334.

Polarity chron	Age (Ma)	Hole U1334A				Measurement type	Hole U1334B				Measurement type	Hole U1334C			
		Range CSF (m)	Best estimate CSF (m)	Best estimate core, section, interval (cm)	Measurement type		Range CSF (m)	Best estimate CSF (m)	Best estimate core, section, interval (cm)	Measurement type		Range CSF (m)	Best estimate CSF (m)	Best estimate core, section, interval (cm)	Measurement type
C1n	0.000	0.000-0.000	0.000	Mudline				Not recovered				Not recovered			
C1n/C1r.1r	0.781	0.700-0.900	0.800	1H-1, 80	Split core			Not identified				Not identified			
C1r.1r/C1r.1n	0.988	1.100-1.100	1.100	1H-1, 110	Split core			Not identified				Not identified			
C1r.1n/C1r.2r	1.072	1.150-1.200	1.175	1H-1, 117.5	Split core			Not identified				Not identified			
C1r.2r/C2n	1.778	1.850-1.900	1.875	1H-2, 37.5	Split core			Not identified				Not identified			
C2n/C2r.1r	1.945	2.000-2.050	2.025	1H-2, 52.5	Split core			Not identified				Not identified			
C2r.1r/C2r.1n	2.128			Not identified				Not identified				Not identified			
C2r.1n/C2r.2r	2.148			Not identified				Not identified				Not identified			
C2r.2r/C2An.1n	2.581			Interpretation uncertain				Not identified				Not identified			
C2An.1n/C2An.1r	3.032			Interpretation uncertain				Not identified				Not identified			
C2An.1r/C2An.2n	3.116			Interpretation uncertain				Not identified				Not identified			
C2An.2n/C2An.2r	3.207			Interpretation uncertain				Not identified				Not identified			
C2An.2r/C2An.3n	3.330			Interpretation uncertain				Not identified				Not identified			
C2An.3n/C2Ar	3.596			Interpretation uncertain				Not identified				Not identified			
C2Ar/C3n.1n	4.187			Interpretation uncertain				Not identified				Not identified			
C3n.1n/C3n.1r	4.300			Interpretation uncertain				Not identified				Not identified			
C3n.1r/C3n.2n	4.493			Interpretation uncertain				Not identified				Not identified			
C3n.2n/C3n.2r	4.631			Interpretation uncertain				Not identified				Not identified			
C3n.2r/C3n.3n	4.799			Interpretation uncertain				Not identified				Not identified			
C3n.3n/C3n.3r	4.896			Interpretation uncertain				Not identified				Not identified			
C3n.3r/C3n.4n	4.997			Interpretation uncertain				Not identified				Not identified			
C3n.4n/C3r	5.235	4.050-4.150	4.100	1H-3, 110	Split core			Not identified				Not identified			
C3r/C3An.1n	6.033	5.250-5.300	5.275	1H-4, 77.5	Split core			Not identified		1.800-1.825	1.813	1H-2, 30-32.5		Split core	
C3An.1n/C3An.1r	6.252	5.700-5.750	5.725	1H-4, 122.5	Split core			Not identified		2.025-2.050	2.038	1H-2, 52.5-55		Split core	
C3An.1r/C3An.2n	6.436	5.900-6.100	6.000	Between Sections 1H-4 and 5	Split core			Not identified		2.200-2.200	2.200	1H-2, 70		Split core	
C3An.2n/C3Ar	6.733	6.300-6.350	6.325	1H-5, 32.5	Split core			Not identified		2.575-2.600	2.588	1H-2, 107.5-110		Split core	
C3Ar/C3Bn	7.140	6.850-6.900	6.875	1H-5, 87.5	Split core			Not identified		3.275-3.275	3.275	1H-3, 27.5		Split core	
C3Bn/C3Br.1r	7.212	7.150-7.250	7.200	1H-5, 120	Split core			Not identified		3.550-3.575	3.563	1H-3, 55-57.5		Split core	
C3Br.1r/C3Br.1n	7.251			Interpretation uncertain				Not identified		3.625-3.625	3.625	1H-3, 62.5		Split core	
C3Br.1n/C3Br.2r	7.285			Interpretation uncertain				Not identified		3.725-3.750	3.738	1H-3, 72.5-75		Split core	
C3Br.2r/C3Br.2n	7.454	7.600-7.650	7.625	1H-6, 12.5	Split core			Not identified		3.975-4.000	3.988	1H-3, 97.5-100		Split core	
C3Br.2n/C3Br.3r	7.489	7.700-7.800	7.750	1H-6, 27.5	Split core			Not identified		4.000-4.025	4.013	1H-3, 100-102.5		Split core	
C3Br.3r/C4n.1n	7.528	7.850-7.900	7.875	1H-6, 37.5	Split core	3.850-3.900	3.875	1H-1, 17.5	Split core	4.025-4.050	4.038	1H-3, 102.5-105		Split core	
C4n.1n/C4n.1r	7.642			Not identified		3.950-4.000	3.975	1H-1, 27.5	Split core	4.300-4.300	4.300	1H-3, 103		Split core	
C4n.1r/C4n.2n	7.695			Not identified		4.000-4.150	4.075	1H-1, 42.5	Split core	4.375-4.400	4.388	1H-3, 137.5-140		Split core	
C4n.2n/C4r.1r	8.108			Not identified		4.600-4.850	4.725	1H-1, 102.5	Split core	5.050-5.075	5.063	1H-4, 55-57.5		Split core	
C4r.1r/C4r.1n	8.254			Not identified		5.000-5.050	5.025	1H-1, 132.5	Split core	5.375-5.375	5.375	1H-4, 87.5		Split core	
C4r.1n/C4r.2r	8.300	8.300-8.300	8.300	2H-1, 10 (?)	Split core	5.100-5.200	5.150	Between Sections 1H-1 and -2 (?)	Split core	5.475-5.575	5.525	1H-4, 97.5-107.5		Split core	
C4r.2r/C4An	8.769	9.300-9.350	9.325	2H-1, 112.5 (?)	Split core	6.250-6.300	6.275	1H-2, 107.5	Split core	6.575-6.600	6.588	1H-5, 57.5-60		Split core	
C4An/C4Ar.1r	9.098	10.100-10.150	10.125	2H-2, 42.5 (?)	Split core	7.150-7.200	7.175	1H-3, 47.5	Split core	7.325-7.425	7.375	1H-5, 132.5-142.5		Split core	
C4Ar.1r/C4Ar.1n	9.312	10.450-10.500	10.475	2H-2, 77.5 (?)	Split core			Interpretation uncertain		7.650-7.675	7.663	1H-6, 15-17.5		Split core	
C4Ar.1n/C4Ar.2r	9.409	10.750-10.750	10.750	2H-2, 105 (?)	Split core	7.900-7.950	7.925	1H-3, 122.5	Split core	7.950-7.975	7.963	1H-6, 45-47.5		Split core	
C4Ar.2r/C4Ar.2n	9.656			Interpretation uncertain		8.300-8.350	8.325	1H-4, 12.5	Split core	8.350-8.375	8.363	1H-6, 45-47.5		Split core	
C4Ar.2n/C4Ar.3r	9.717			Interpretation uncertain		8.500-8.550	8.525	1H-4, 32.5	Split core			Not identified			
C4Ar.3r/C5n.1n	9.779			Interpretation uncertain		8.600-8.650	8.625	1H-4, 42.5	Split core			Not identified			
C5n.1n/C5n.1r	9.934			Interpretation uncertain				Interpretation uncertain				Not identified			
C5n.1r/C5n.2n	9.987			Interpretation uncertain				Interpretation uncertain				Not identified			
C5n.2n/C5r.1r	11.040	13.400-13.450	13.425	2H-4, 72.5	Split core	10.850-10.950	10.900	1H-5, 120	Split core			Not identified			
C5r.1r/C5r.1n	11.118	13.500-13.600	13.550	2H-4, 85	Split core	11.000-11.050	11.025	1H-5, 132.5	Split core			Not identified			
C5r.1n/C5r.2r	11.154	13.650-13.750	13.700	2H-4, 95	Split core	11.100-11.300	11.200	Between Sections 1H-5 and 6	Split core			Not identified			
Excursion? (top)				Not identified		11.650-11.650	11.650	1H-6, 57.5	Split core	10.560-10.560	10.560	2H-2, 10		Split core	
Excursion? (bottom)				Not identified		11.750-11.800	11.775	1H-6, 45	Split core	10.660-10.710	10.685	2H-2, 22.5		Split core	
C5r.2r/C5r.2n	11.554	14.650-14.700	14.675	2H-5, 47.5	Split core	12.200-12.250	12.225	1H-6, 102.5	Split core	11.060-11.160	11.110	2H-2, 65		Split core	
C5r.2n/C5r.3r	11.614	14.800-14.900	14.850	2H-5, 65	Split core	12.450-12.500	12.475	1H-6, 127.5	Split core	11.260-11.410	11.335	2H-2, 85		Split core	
C5r.3r/C5An.1n	12.014	15.900-15.950	15.925	2H-6, 22.5	Split core			Poor quality paleomagnetic data		12.410-12.460	12.435	2H-3, 45-50		Split core	
C5An.1n/C5An.1r	12.116	16.200-16.250	16.225	2H-6, 52.5	Split core			Poor quality paleomagnetic data		12.760-12.860	12.810	2H-3, 85		Split core	
C5An.1r/C5An.2n	12.207	16.450-16.600	16.525	2H-6, 82.5	Split core			Poor quality paleomagnetic data		13.010-13.160	13.085	2H-3, 105-120		Split core	
C5An.2n/C5Ar.1r	12.415	17.400-17.500	17.450	2H-7, 25	Split core			Poor quality paleomagnetic data		14.060-14.060	14.060	2H-4, 30		Split core	
C5Ar.1r/C5Ar.1n	12.730	18.250-18.350	18.300	3H-1, 60	Split core			Poor quality paleomagnetic data		15.810-15.910	15.860	2H-5, 90		Split core	
C5Ar.1n/C5Ar.2r	12.765	18.350-18.450	18.400	3H-1, 70	Split core			Poor quality paleomagnetic data		16.010-16.110	16.060	2H-5, 110		Split core	
C5Ar.2r/C5Ar.2n	12.820	18.800-18.850	18.825	3H-1, 112.5	Split core			Poor quality paleomagnetic data		16.310-16.360	16.335	2H-5, 137.5		Split core	
C5Ar.2n/C5Ar.3r	12.878	18.900-19.000	18.950	3H-1, 125	Split core			Poor quality paleomagnetic data		16.560-16.610	16.585	2H-6, 12.5		Split core	
C5Ar.3r/C5AAn	13.015	19.600-19.750	19.675	3H-2, 47.5	Split core			Poor quality paleomagnetic data		17.210-17.260	17.235	2H-6, 77.5		Split core	
C5AAn/C5AAn	13.183	20.300-20.350	20.325	3H-2, 112.5	Split core	17.500-17.600	17.550	2H-3, 135	Split core	18.060-18.160	18.110	2H-7, 15		Split core	
C5AAn/C5ABn	13.369	21.050-21.300	21.175	3H-3, 47.5	Split core	18.350-18.400	18.375	2H-4, 67.5	Split core			Not identified (in gap between cores)			
C5ABn/C5ABr	13.605	22.650-22.750	22.700	3H-4, 50	Split core	20.000-20.100	20.050	2H-5, 85	Split core			Not identified (in gap between cores)			
C5ABr/C5ACn	13.734	23.350-23.450	23.400	3H-4, 120	Split core	20.800-20.900	20.850	2H-6, 15	Split core			Not identified (in gap between cores)			
C5ACn/C5ACr	14.095	24.700-24.800	24.750	3H-5, 105	Split core	22.100-22.300	22.200	Between Sections 2H-6 and 7	Split core	20.000-20.100	20.050	3H-2, 25		Split core	
C5ACr/C5ADn	14.194	24.950-25.050	25.000	3H-5, 130	Split core	22.400-22.500	22.450	2H-7, 25	Split core	20.250-20.350	20.300	3H-2, 50		Split core	
C5ADn/C5ADr	14.581	26.600-26.850	26.725	Between Sections 3H-6 and 7	Split core	24.000-24.100	24.050	3H-1, 135	Split core	22.050-22.100	22.075	3H-3, 77.5		Split core	
Excursion? (top)				Not identified				Not identified		22.400-22.450	22.425	3H-3, 112.5		Split core	
Excursion? (bottom)				Not identified				Not identified		22.450-22.500	22.475	3H-3, 117.5		Split core	
C5ADr/C5Bn.1n	14.784			Not identified (in gap between cores)		24.900-25.000	24.950	3H-2, 75	Split core	23.000-23.000	23.000	3H-4, 20		Split core	
C5Bn.1n/C5Bn.1r	14.877			Not identified (in gap between cores)		25.250-25.350	25.300	3H-2, 110	Split core	23.300-23.350	23.325	3H-4, 52.5		Split core	
C5Bn.1r/C5Bn.2n	15.032			Not identified (in gap between cores)		26.050-26.100	26.075	3H-3, 37.5	Split core	24.050-24.150	24.100	3H-4, 130		Split core	
C5Bn.2n/C5Br	15.160	27.700-27.900	27.800	4H-1, 60	Split core	26.800-26.900	26.850	3H-3, 115	Split core	24.650-24.800	24.725	3H-5, 42.5		Split core	
Excursion? (top)		28.500-28.500	28.500	4H-1, 130	Split core	27.550-27.550	27.550	3H-4, 35	Split core	25.350-25.400	25.375	3H-5, 107.5		Split core	
Excursion? (bottom)		28.550-28.600	28.575	4H-1, 137.5	Split core	27.650-27.650	27.650	3H-4, 45	Split core	25.450-25.500	25.475	3H-5, 117.5		Split core	
C5Br/C5Cn.1n	15.974	33.100-33.300	33.200	Between Sections 4H-4 and 5	Split core	31.450-31.550	31.500	3H-6, 130	Split core			Not identified (in gap between cores)			
C5Cn.1n/C5Cn.1r	16.268	34.600-34.800	34.700	Between Sections 4H-5 and 6	Split core			Not identified (in gap between cores)				Not identified (in gap between cores)			
C5Cn.1r/C5Cn.2n	16.303	35.050-35.100	35.075	4H-6, 37.5	Split core			Not identified (in gap between cores)		28.950-29.000	28.975	4H-1, 47.5		Split core	
C5Cn.2n/C5Cn.2r	16.472	35.350-35.400	35.375	4H-6, 67.5	Split core	32.600-32.600	32.600	4H-1, 40	Split core	29.150-29.200	29.175	4H-1, 67.5		Split core	
C5Cn.2r/C5Cn.3n	16.543	35.750-35.850	35.800	4H-6, 110	Split core	33.000-33.100	33.050	4H-1, 85	Split core	29.500-29.550	29.525	4H-1, 102.5		Split core	
C5Cn.3n/C5Cr	16.721	36.050-36.300	36.175	Between Sections 4H-6 and 7	Split core	33.400-33.450	33.425	4H-1, 122.5	Split core	29.850-29.900	29.875	4H-1, 137.5			