Additional file 9: Shared Haplotypes between East Timor and 16 surrounding populations based on HVS-I data (reading frame: 16080-16180 16195-16354)

Population	East Timor	Nusa Tenggara		a Sumatra	Mentawai		Peninsular Malaysia										China (Mixed Han)				Admiralty Islands		Polynesia	Australia (Indi
Reference	[3,this study]	[3,5,24,65]	[5,24] [5,2			[5,41]	[17,57]	[24]	[5,24]	[24,65]	[24,28]	[58]	[59]	[60]	[62]	[24,45]	[63]	[61]	[66]	[34,64,67]	[67,68]	[29]	[65,67]	[34,69
number of samples	362	1699	570 97			499	470	157	437	74	483	187	214	190	293	718	262	692	227		203	703	394	146
number of haplotypes	94	345	138 49	107	16	61	158	102	153	36	140	119	127	108	99	86	194	306	74	72	41	102	56	50
ET030	39	33	14 2	6	0	1	2	10	61	3	43	0	0	0	0	24	1	1	0	0	0	0	0	0
ET067	26	82	11 6	15	7	22	11	3	32	4	37	3	1	1	9	22	3	3	0	0	2	10	1	0
ET045	24	88	0 0		0	0	1	3	13	0	12	1	0	0	3	6	0	0	0	0	0	0	0	0
ET012	18	53	17 2	8	0	7	3	5	22	10	23	2	6	1	1	40	10	69	2	3	0	6	0	7
ET152	15	14	2 0	1	0	0	0	3	1	0	14	0	0	0	0	14	1	0	0	0	0	0	0	0
ET165	13	7	0 0	0	0	0	0	2	9	6	0	0	0	0	0	0	0	0	0	3	70	305	225	0
ET131	12	72	11 2	2	0	0	1	2	7	8	15	1	0	0	0	2	0	2	0	0	0	0	0	(
ET100	11	15	0 0	0	0	0	0	0	2	4	0	0	0	0	0	0	0	0	31	33	27	5	1	0
ET161	11	36	3 0		0	36	1	7	18	3	40	0	0	0	2	41	2	3	0		16	77	41	(
ET202	8	56	4 0		4	3	2	1	16	0	3	0	0	0	0	1	0	0	0	0	2	6	0	0
ET187	7	59	0 0		0	0	<u></u>	1	8	2	20	0	2	0	15	47	0	2	0	0	0	3	5	0
ET319	7	19	36 2		0	13	2	2	22	1	15	0	0	1	1	12	0	1	0	0	0	0	0	0
ET212	6	56	5 0		23	21	3	2	7	1	24	0	0	0	0	0	0	5	0	0	3	1	0	
ET103	5	1	0 0		0	5	1	0	0	0	0	1	1	0	0	1	1	1	0	0	0	1	2	
ET103	5	11	3 0		0	0	1	1	1	1	2	0	0	0	0	0	0	0	0	0	1	2	0	
ET065	4	0	0 0		0	0	0	0	0	0	0	0	0	0	0	7	0	0	0	0	0	0	0	
ET272	4	3	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ET017	3	3	0 0		0	0	0	0	0	2	0	0	0	0	0	0	0	0	8	4	2	5	0	
ET071	3	0	10 0		0	0	0	0	0	0	3	1	1	0	1	0	0	0	0	0	0	0	0	
ET074	3	33	0 0		0	12	1	1	8	0	1	0	0	0	0	9	0	0	0	0	0	0	0	
ET186	3	0	0 0		0	0	0	0	0	0	2	0	0	0	3	0	0	0	0	0	0	0	0	
ET234	2	10	0 0		0	0	0	1	0	1	0	0	0	0	0	0	0	0	13	1	0	0	0	
ET274	3	0	0 0		0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ET008	3	1	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ET008	2	54			-		3			-	3	- 0		1			3				-	_	0	(
ET021 ET094	2	22	5 4		0	1	0	4	5	0	0	0	2		3	0	0	15	0	0	0	0	0	
		1	0 0		0	0		0		0	0		0	0	0	0	•	0	0	0	0	0	- J	
ET153	2	6	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	7	0	0	0	0	
ET209	2	6	0 0		-	0	0		3	0	0	0	0	3		-	0	0	0	Ū			0	(
ET278	2	5		9	0	1	3	4				0	3		0	0	0	0	0	0	0	0	0	
FJ838905 [3]	2	10	1 6		0	0	3	0	1	0	1	2	1	8	8	0	3	1	0	3	0	0	0	
ET015		0	0 0		0	0	0	0	1	0	- 0	0	0		0	0	0	1	0	0	0	0	0	,
ET037	1	16	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	2	0	0	0	
ET077	1	0	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ET078	1	26	1 5		0	8	24	0	3	1	0	8	7	10	3	0	2	0	0	0	0	0	0	
ET079	1	0	0 0		0	0	0	0	0	0	0	0	0	11	0	3	0	0	0	0	0	0	0	
ET110	1	1	0 4		0	0	0	1	0	0	2	0	0	0	1	0	0	0	0	0	0	0	0	
ET122	1	4	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
ET126	1	15	1 0		0	0	11	0	3	0	0	1	0	1	0	0	1	5	0	0	0	0	0	(
ET181	1	7	0 0		0	5	0	0	0	0	3	0	0	0	0	19	0	0	0	0	0	0	0	(
ET194	1	0	0 0		0	1	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1
ET196	1	9	0 0		0	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
ET224	1	0	0 0		0	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	(
ET225	1	0	0 0	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3	5	9	1
ET257	1	7	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
ET259	1	5	0 0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
ET264	1	0	0 0		0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	(
ET277	1	1	1 0		0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	(
ET283	1	0	0 0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	(
FJ838923 [3]	1	1	0 0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
FJ838931 [3]	1	0	0 0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	(
FJ838932 [3]	1	0	0 0		0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	(
umber of shared haplotypes	51	37	17 10		3	14	19	18	25	15	21	10	9	10	12	16	10	14	6	8	9	12	7	
ortion of shared haplotypes [%]	54,26	10,72	12,32 20,4	11 16,82	18,75	22,95	12,03	17,65	16,34	41,67	15,00	8,40	7,09	9,26	12,12	18,60	5,15	4,58	8,11	11,11	21,95	11,76	12,50	4,
viduals with shared haplotypes	268	842	136 34	74	34	136	65	53	247	48	266	21	24	28	50	249	27	110	62	50	126	426	284	1-

one example haplotype given per shared haplotype highest shared proportion marked in green lowest shared proportion marked in red