

Table S2: Runs of homozygosity segments shared by over 50% of Shagya Arabians.

Chr.	Length (MB)	Begin	End	Annotated genes
1	0.02	9501779	9519364	-
1	0.64	107739971	108378967	TARSL2, TJP1, FAM189A1 OPRD1, YTHDF2, GMEB1, TAF12, RAB42, RCC1, PHACTR4, MED18, SESN2, ATP5IF1, DNAJC8, PTAFR, EYA3, XKR8, SMPDL3B, RPA2, THEMIS2, PPP1R8, STX12,
2	1.06	27344713	28404281	FAM76A, IFI6, FGR, AHDC1 NKX1-1, UVSSA, MAEA, CTBP1, SPON2, RNF212, FGFR1L, IDUA, SLC26A1, DGKQ, TMEM175, GAK, MIR9070, CPLX1,
3	0.93	120425335	121351057	PCGF3, SLC49A3, MYL5, ATP5ME, PDE6B, PIGG ZNF142, BCS1L, RNF25, STK36, TTL4, CYP27A1,
6	0.18	8018928	8201303	PRKAG3
7	0.25	40755980	41002946	NTM
7	0.88	41060288	41938373	NTM, OPCML RHOF, SETD1B, HPD, PSMD9, WDR66, BCL7A, MLXIP, IL31, LRRC43, B3GNT4, DIABLO, VPS33A, CLIP1, ZCCHC8,
8	0.60	24634587	25231545	RSRC2, KNTC1 KMT5A, RILPL2, SNRNP35, RILPL1, TMED2, DDX55, EIF2B1, GTF2H3, TCTN2, ATP6V0A2, DNAH10
8	0.27	25825349	26097405	-
8	0.00	61839365	61839365	
9	0.53	31284516	31817479	RB1CC1, ALKAL1
9	0.76	31853320	32616875	ST18, PCMTD1
9	0.22	32677402	32897046	PXDNL
9	0.52	33532618	34053277	SNTG1
9	0.87	40515338	41382074	TRIQK
9	0.58	41416774	41992448	FAM92A, RBM12B, TMEM67
9	0.76	41998238	42758961	TMEM67, PDP1, CDH17, GEM, RAD54B, VIRMA
9	0.03	42822721	42852068	ESRP1, DPY19L4
9	0.01	45565108	45576777	MTDH LAPTM4B, MATN2, RPL30, ERICH5, RIDA, POP1, NIPAL2,
9	0.92	45613807	46532261	KCNS2, STK3
9	0.61	79742728	80350559	FAM135B, COL22A1 KRT10B, KRT28, KRT27, KRT26, KRT25, KRT24, KRT222, SMARCE1, CCR7, TNS4, IGFBP4, TOP2A, GJD3, RARA, CDC6, WIPF2, RAPGEFL1, CASC3, MSL1, NR1D1, THRA, MED24, CSF3, PSMD3, GSDMA, LRRC3C, ORMDL3,
11	0.72	21898983	22614034	GSDMB
11	0.00	22654340	22654340	IKZF3
11	0.05	22703722	22750090	IKZF3, GRB7 CDK12, MED1, FBXL20, STAC2, RPL19, CACNB1, ARL5C,
11	0.31	22880941	23186362	PLXDC1 NOG, DGKE, TRIM25, COIL, SCPEP1, AKAP1, MSI2,
11	1.12	31314776	32434775	CCDC182
11	0.04	32617562	32658713	DYNLL2 LPO, MPO, TSPOAP1, RNF43, SUPT4H1, HSF5, MTMR4,
11	0.58	32803157	33383463	TEX14, RAD51C, PPM1E
11	0.23	34276336	34508997	MED13
11	0.01	34577854	34586693	BRP1
11	0.12	34683002	34804064	BRP1
15	1.04	80216896	81259084	NBAS, FAM84A
16	0.16	33620538	33782347	ERC2
16	0.35	41847777	42194451	LTF, CCRL2, CCR5, CCR2, CCR3
18	0.56	49358178	49917777	SSB, METTL5, UBR3, MYO3B
18	0.51	50413659	50924378	TLK1, METTL8, DCAF17
18	0.09	81688757	81776604	CREB1, METTL21A
19	0.07	34349123	34421059	XXYL1
23	0.42	25249949	25673636	GLIS3, SLC1A1, SPATA6L, CDC37L1, AK3, RCL1