

Table S1: Genes overlapping in at least two domesticated species

GENE NAME	OVERLAPPING SPECIES	ENSEMBL ID	GENE FUNCTION (UniProt)	PATHWAY ANNOTATION	RELATED DISORDERS
ADAMTS13	cattle, cat	ENSG00000160323	Cleaves the vWF multimers in plasma into smaller forms thereby controlling vWF-mediated platelet thrombus formation.	Metabolism of proteins (Reactome)	Thrombotic thrombocytopenic purpura, familial (OMIM)
ATXN7L1	cat, dog	ENSG00000146776	N/A	N/A	N/A
BRAF	cat, horse	ENSG00000157764	Protein kinase involved in the transduction of mitogenic signals from the cell membrane to the nucleus. May play a role in the postsynaptic responses of hippocampal neuron. Phosphorylates MAP2K1, and thereby contributes to the MAP kinase signal transduction pathway.	EGFR tyrosine kinase inhibitor resistance, Endocrine resistance, MAPK signaling pathway, ErbB signaling pathway, Rap1 signaling pathway, cAMP signaling pathway, Chemokine signaling pathway, FoxO signaling pathway, mTOR signaling pathway, Vascular smooth muscle contraction, Dorso-ventral axis formation, Focal adhesion, Natural killer cell mediated cytotoxicity, Long-term potentiation, Neurotrophin signaling pathway, Serotonergic synapse, Long-term depression, Regulation of actin cytoskeleton, Insulin signaling pathway, Progesterone-mediated oocyte maturation, Alcoholism, Hepatitis C, Pathways in cancer, Proteoglycans in cancer, Colorectal cancer, Renal cell carcinoma, Pancreatic cancer, Endometrial cancer, Glioma, Prostate cancer, Thyroid cancer, Melanoma, Bladder cancer, Chronic myeloid leukemia, Acute myeloid leukemia, Non-small cell lung cancer, Breast cancer (KEGG); VEGF signaling pathway, T cell activation, Interleukin signaling pathway, Inflammation mediated by chemokine and cytokine signaling pathway, Integrin signaling pathway, EGF receptor signaling pathway, Ras Pathway, Angiogenesis, CKKR signaling map, PDGF signaling pathway, B cell activation (PANTHER).	Thyroid cancer, Malignant melanoma, Noonan syndrome and related disorders, Langerhans cell histiocytosis (KEGG); Adenocarcinoma of lung, somatic; Cardiofaciocutaneous syndrome; Colorectal cancer, somatic; LEOPARD syndrome; Melanoma, malignant, somatic; Non-small cell lung cancer, somatic; Noonan syndrome (OMIM)
CLEC5A	dog, cat	ENSG00000258227	Functions as a positive regulator of osteoclastogenesis. Cell surface receptor that signals via TYROBP. Regulates inflammatory responses. Acts as a key regulator of synovial injury and bone erosion during autoimmune joint inflammation (By similarity). Critical macrophage receptor for dengue virus serotypes 1-4.	Immune System (Reactome)	N/A
DCC	horse, cat	ENSG00000187323	Receptor for netrin required for axon guidance. Mediates axon attraction of neuronal growth cones in the developing nervous system upon ligand binding. Its association with UNC5 proteins may trigger signaling for axon repulsion. It also acts as a dependence receptor required for apoptosis induction when not associated with netrin ligand. Implicated as a tumor suppressor gene.	Axon guidance mediated by netrin, Axon guidance mediated by Slii/Robo (PANTHER)	Colorectal cancer, somatic; Esophageal carcinoma, somatic; Mirror movements (OMIM)
FAM172A	dog, cattle	ENSG00000113391	N/A	N/A	N/A
GRIK3	dog, cattle	ENSG00000163873	Receptor for glutamate that functions as ligand-gated ion channel in the central nervous system and plays an important role in excitatory synaptic transmission. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. The postsynaptic actions of Glu are mediated by a variety of receptors that are named according to their selective agonists.	Neuroactive ligand-receptor interaction; Glutamatergic synapse (KEGG); Huntington disease, Ionotropic glutamate receptor pathway, Metabotropic glutamate receptor group III pathway (PANTHER)	schizophrenia, obsessive-compulsive disorder, alcohol dependence (MalaCards)
NRG2	dog, cat, cattle	ENSG00000158458	Direct ligand for ERBB3 and ERBB4 tyrosine kinase receptors. Concomitantly recruits ERBB1 and ERBB2 co-receptors, resulting in ligand-stimulated tyrosine phosphorylation and activation of the ERBB receptors. May also promote the heterodimerization with the EGF receptor.	EGF receptor signaling pathway (PANTHER), EGFR tyrosine kinase inhibitor resistance, ErbB signaling pathway (KEGG).	Hereditary motor and sensory neuropathy (Charcot-Marie-Tooth disease) (MalaCards)
PLAC8L1	cattle, cat	ENSG00000173261	N/A	N/A	N/A
RNPC3	dog, cat	ENSG00000185946	Participates in pre-mRNA U12-dependent splicing, performed by the minor spliceosome which removes U12-type introns. U12-type introns comprises less than 1% of all non-coding sequences. Binds to the 3'-stem-loop of m7G-capped U12 snRNA.	Gene Expression (Reactome)	growth hormone deficiency, parainfluenza virus type 3, idiopathic inflammatory myopathy, rheumatic disease (MalaCards)
SEC24A	cat, horse	ENSG00000113615	Component of the COPII coat, that covers ER-derived vesicles involved in transport from the endoplasmic reticulum to the Golgi apparatus. COPII acts in the cytoplasm to promote the transport of secretory, plasma membrane, and vacuolar proteins from the endoplasmic reticulum to the Golgi complex.	Protein processing in endoplasmic reticulum (KEGG)	N/A
SMG6	cat, horse	ENSG00000070366	Component of the telomerase ribonucleoprotein (RNP) complex that is essential for the replication of chromosome termini. May have a general role in telomere regulation. Promotes in vitro the ability of TERT to elongate telomeres. Overexpression induces telomere uncapping, chromosomal end-to-end fusions (telomeric DNA persists at the fusion points) and did not perturb TRF2 telomeric localization. Binds to the single-stranded 5'-(GTGTGG)4GTGT-3' telomeric DNA, but not to a telomerase RNA template component (TER). Plays a role in nonsense-mediated mRNA decay. Is thought to provide a link to the mRNA degradation machinery as it has endonuclease activity required to initiate NMD, and to serve as an adapter for UPF1 to protein phosphatase 2A (PP2A), thereby triggering UPF1 dephosphorylation. Degrades single-stranded RNA (ssRNA), but not dsDNA or dsRNA.	mRNA surveillance pathway (KEGG)	tricuspid valve insufficiency (MalaCards)
STK10	dog, cattle	ENSG00000072786	Serine/threonine-protein kinase involved in regulation of lymphocyte migration. Phosphorylates MSN, and possibly PLK1. Involved in regulation of lymphocyte migration by mediating phosphorylation of ERM proteins such as MSN. Acts as a negative regulator of MAP3K1/MEK1. May also act as a cell cycle regulator by acting as a polo kinase kinase; mediates phosphorylation of PLK1 in vitro; however such data require additional evidences in vivo. (UniProt)	N/A	Testicular germ cell tumor, seminal vesicle tumor, ocal disease, anus disease, biliary dyskinesia (MalaCards)
TMEM132D	dog, cattle	ENSG00000151952	May serve as a cell-surface marker for oligodendrocyte differentiation.	N/A	pithrus pubis infestation, lice infestation, parasitic ectoparasitic infectious disease (MalaCards)
VEZT	dog, cat	ENSG00000028203	Plays a pivotal role in the establishment of adherens junctions and their maintenance in adult life. In case of <i>Listeria</i> infection, promotes bacterial internalization by participating in myosin VIIa recruitment to the entry site.	N/A	N/A

Table S2: Overlapping genes in AMH and domesticated animals

GENE NAME	OVERLAPPING SPECIES	ENSEMBL ID	GENE FUNCTION (UniProt)	PATHWAY ANNOTATION	RELATED DISORDERS
AMBRA1	horse	ENSG00000110497	Regulates autophagy and development of the nervous system. Involved in autophagy in controlling protein turnover during neuronal development, and in regulating normal cell survival and proliferation (By similarity).	Cellular responses to stress (Reactome)	N/A
BRAF	cat, horse	ENSG00000157764	Protein kinase involved in the transduction of mitogenic signals from the cell membrane to the nucleus. May play a role in the postsynaptic responses of hippocampal neuron. Phosphorylates MAP2K1, and thereby contributes to the MAP kinase signal transduction pathway.	EGFR tyrosine kinase inhibitor resistance, Endocrine resistance, MAPK signaling pathway, ERB signaling pathway, Rap1 signaling pathway, cAMP signaling pathway, Chemokine signaling pathway, FoxO signaling pathway, mTOR signaling pathway, Vascular smooth muscle contraction, Dorso-ventral axis formation, Focal adhesion, Natural killer cell mediated cytotoxicity, Long-term potentiation, Neurotrophin signaling pathway, Serotonergic synapse, Long-term depression, Regulation of actin cytoskeleton, Insulin signaling pathway, Progesterone-mediated oocyte maturation, Alcoholism, Hepatitis C, Pathways in cancer, Proteoglycans in cancer, Colorectal cancer, Renal cell carcinoma, Pancreatic cancer, Endometrial cancer, Glioma, Prostate cancer, Thyroid cancer, Melanoma, Bladder cancer, Chronic myeloid leukemia, Acute myeloid leukemia, Non-small cell lung cancer, Breast cancer (KEGG); VEGF signaling pathway, T cell activation, Interleukin signaling pathway, Inflammation mediated by chemokine and cytokine signaling pathway, Integrin signaling pathway, EGF receptor signaling pathway, Ras Pathway, Angiogenesis, CCKR signaling map, PDGF signaling pathway, B cell activation (PANTHER)	Thyroid cancer; Malignant melanoma; Noonan syndrome and related disorders; Langerhans cell histiocytosis (KEGG); Adenocarcinoma of lung, somatic; Cardiovascular disease; Colorectal cancer, somatic; LEOPARD syndrome; Melanoma, malignant, somatic; Non-small cell lung cancer, somatic; Noonan syndrome (OMIM)
CACNA1D	horse	ENSG00000157388	Voltage-sensitive calcium channels (VSCC) mediate the entry of calcium ions into excitable cells and are also involved in a variety of calcium-dependent processes, including muscle contraction, hormone or neurotransmitter release, gene expression, cell motility, cell division and cell death. The isoform alpha-1D gives rise to L-type calcium currents. Long-lasting (L-type) calcium channels belong to the "high-voltage activated" (HVA) group. They are blocked by dihydropyridines (DHP), phenylalkylamines, benzothiazepines, and by omega-agatoxin-IIIa (omega-Aga-IIIa). They are however insensitive to omega-conotoxin-GVIA (omega-Ctx-GVIA) and omega-agatoxin-IVA (omega-Aga-IVA).	Amphetamine addiction, Vascular smooth muscle contraction, MAPK signaling pathway, Calcium signaling pathway, cGMP-PKG signaling pathway, cAMP signaling pathway, Cardiac muscle contraction, Adrenergic signaling in cardiomyocytes, Tight junction, Circadian entrainment, Retrograde endocannabinoid signaling, Glutamatergic synapse, Cholinergic synapse, Serotonergic synapse, GABAergic synapse, Dopaminergic synapse, Insulin secretion, GnRH signaling pathway, Oxytocin signaling pathway, Renin secretion, Aldosterone synthesis and secretion, Type II diabetes mellitus, Carbohydrate digestion and absorption, Alzheimer's disease, Hypertrophic cardiomyopathy, Arrhythmogenic right ventricular cardiomyopathy, Dilated cardiomyopathy (KEGG); Oxytocin receptor mediated signaling pathway, Nicotinic acetylcholine receptor signaling pathway, Beta2 adrenergic receptor signaling pathway, SH2 type receptor mediated signaling pathway, Gonadotropin-releasing hormone receptor pathway, Alzheimer disease-amyloid secretase pathway, Beta1 adrenergic receptor signaling pathway (PANTHER)	Primary aldosteronism, seizures, and neurologic abnormalities; Sinusoidal node dysfunction and deafness (OMIM)
COA5	dog	ENSG00000183513	Involved in an early step of the mitochondrial complex IV assembly process.	N/A	Cardioencephalomyopathy, fatal infantile, due to cytochrome c oxidase deficiency 3 (OMIM), Cytochrome c oxidase (COX) deficiency (KEGG)
COL11A1	dog	ENSG000000060718	May play an important role in fibrillogenesis by controlling lateral growth of collagen II fibrils.	Integrin signaling pathway (PANTHER); Extracellular matrix organization (Reactome)	Fibrochondrogenesis 1; Marshall syndrome; Stickler syndrome, type II; Lumbar disc herniation, susceptibility to (OMIM)
COQ10B	dog	ENSG00000115520	Required for the function of coenzyme Q in the respiratory chain. May serve as a chaperone or may be involved in the transport of Q6 from its site of synthesis to the catalytic sites of the respiratory complexes (By similarity).	Metabolism (Reactome)	N/A
DLGAP1	horse	ENSG00000170579	Part of the postsynaptic scaffold in neuronal cells.	Glutamatergic synapse (KEGG)	N/A
ERBB4	cattle	ENSG00000178568	Tyrosine-protein kinase that plays an essential role as cell surface receptor for neuregulins and EGF family members and regulates development of the heart, the central nervous system and the mammary gland, gene transcription, cell proliferation, differentiation, migration and apoptosis. Required for normal cardiac muscle differentiation during embryonic development, and for postnatal cardiomyocyte proliferation. Required for normal development of the embryonic central nervous system, especially for normal neural crest cell migration and normal axon guidance. Required for mammary gland differentiation, induction of milk proteins and lactation. Acts as cell-surface receptor for the neuregulins NRG1, NRG2, NRG3 and NRG4 and the EGF family members ETO, ERBB and HBEGF. Ligand binding triggers receptor dimerization and autophosphorylation at specific tyrosine residues that then serve as binding sites for scaffold proteins and effectors. Ligand specificity and signaling is modulated by alternative splicing, proteolytic processing and by the formation of heterodimers with other ERBB family members, thereby creating multiple combinations of intracellular phosphotyrosines that trigger ligand- and context-specific cellular responses. Mediates phosphorylation of SHC1 and activation of the MAP kinases MAPK1/ERK2 and MAPK3/ERK1. Isoform JM-A CYT-1 and isoform JM-B CYT-1 phosphorylate PIK3R1, leading to the activation of phosphatidylinositol 3-kinase and AKT1 and protect cells against apoptosis. Isoform JM-A CYT-1 and isoform JM-B CYT-1 mediate reorganization of the actin cytoskeleton and promote cell migration in response to NRG1. Isoform JM-A CYT-2 and isoform JM-B CYT-2 lack the phosphotyrosine that mediates interaction with PIK3R1, and hence do not phosphorylate PIK3R1, do not protect cells against apoptosis, and do not promote reorganization of the actin cytoskeleton and cell migration. Proteolytic processing of isoform JM-A CYT-1 and isoform JM-A CYT-2 gives rise to the corresponding soluble intracellular domains (iCD) that translocate to the nucleus, promote nuclear import of STAT5A, activation of STAT5A, mammary epithelium differentiation, cell proliferation and activation of gene expression. The ERBB4 soluble intracellular domains (iCD) colocalize with	Erbb signaling pathway, Calcium signaling pathway, Endocytosis, Proteoglycans in cancer (KEGG); Alzheimer disease-prenilin pathway, EGF receptor signaling pathway, Cadherin signaling pathway (PANTHER)	Amyotrophic lateral sclerosis 19 (OMIM)
FAM172A	cattle, dog	ENSG00000113391	N/A	N/A	N/A
GGT7	dog	ENSG00000131067	Cleaves glutathione conjugates.	N/A	N/A
GRIA1	cat	ENSG00000155511	Ionotropic glutamate receptor. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. Binding of the excitatory neurotransmitter L-glutamate induces a conformation change, leading to the opening of the cation channel, and thereby converts the chemical signal to an electrical impulse. The receptor then desensitizes rapidly and enters a transient inactive state, characterized by the presence of bound agonist. In the presence of CACNG4 or CACNG7 or CACNG8, shows resensitization which is characterized by a delayed accumulation of current flux upon continued stimulation of the channel.	Amphetamine addiction, Circadian entrainment, Long-term depression, Nicotine addiction, cAMP signaling pathway, Neuroactive ligand-receptor interaction, Long-term potentiation, Retrograde endocannabinoid signaling, Glutamatergic synapse, Dopaminergic synapse, Amyotrophic lateral sclerosis (ALS) (KEGG)	status epilepticus; fragile x syndrome; limbic encephalitis; schizoprenia; amyotrophic lateral sclerosis 1 (MalaCards)
GRIK3	dog, cattle	ENSG00000163873	Receptor for glutamate that functions as ligand-gated ion channel in the central nervous system and plays an important role in excitatory synaptic transmission. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. The postsynaptic actions of Glu are mediated by a variety of receptors that are named according to their selective agonists. This receptor binds domoate > kainate >> L-glutamate = quisqualate >> AMPA = NMDA.	Glutamatergic synapse, Neuroactive ligand-receptor interaction (KEGG); Huntington disease (PANTHER)	schizophrenia; alcohol dependence (MalaCards)
HSD3B7	cat	ENSG00000099377	The 3-beta-HSD enzymatic system plays a crucial role in the biosynthesis of all classes of hormonal steroids. HSD VII is active against four 7-alpha-hydroxylated steroids. Does not metabolize several different C(19/21) steroids as substrates. Involved in bile acid synthesis (PubMed:11067870). Plays a key role in cell positioning and movement in lymphoid tissues by mediating degradation of 7-alpha,25-dihydroxycholesterol (7-alpha,25-OHC); 7-alpha,25-OHC acts as a ligand for the G protein-coupled receptor GPR183/EBI2, a chemotactic receptor for a number of lymphoid cells.	Primary bile acid biosynthesis, Metabolic pathways (KEGG); Androgen/estrogen/progesterone biosynthesis (PANTHER)	Bile acid synthesis defect, congenital, 1 (OMIM)
HSPD1	dog	ENSG00000144381	Implicated in mitochondrial protein import and macromolecular assembly. May facilitate the correct folding of imported proteins. May also prevent misfolding and promote the refolding and proper assembly of unfolded polypeptides generated under stress conditions in the mitochondrial matrix.	Gene Expression, Metabolism of proteins (Reactome)	Leukodystrophy, hypomyelinating, 4; Spastic paraplegia 13, autosomal dominant (OMIM)
HSPE1	dog	ENSG00000115541	Eukaryotic CPN10 homolog which is essential for mitochondrial protein biogenesis, together with CPN60. Binds to CPN60 in the presence of Mg-ATP and suppresses the ATPase activity of the latter.	N/A	exocervical carcinoma, senneelsu fever; Leprosy; carion's disease (MalaCards)
ITGA9	cat	ENSG00000144688	Integrin alpha-9beta-1 (ITGA9-ITGB1) is a receptor for VCAM1, cytostatin and osteopontin. It recognizes the sequence A-E-I-D-G-I-E-L in cytostatin.	Cell adhesion molecules, PI3K-Akt signaling, Focal adhesion, ECM-receptor interaction, Regulation of actin cytoskeleton, Hypertrophic cardiomyopathy, Arrhythmogenic right ventricular cardiomyopathy, Dilated cardiomyopathy (KEGG); Integrin signaling pathway, Inflammation mediated by chemokine and cytokine signaling pathway (PANTHER)	congenital muscular dystrophy with hyperlaxity, multi-torse syndrome; lynch syndrome (MalaCards)
LRP1B	cattle	ENSG00000168702	cellular cell surface proteins that bind and internalize ligands in the process of receptor-mediated endocytosis.	Alzheimer disease-prenilin pathway (PANTHER)	endocervical carcinoma (MalaCards)
LYST	dog	ENSG00000143669	May be required for sorting endosomal resident proteins into late multivesicular endosomes by a mechanism involving microtubules.	Tropae, piperidine and pyridine alkaloid biosynthesis (KEGG)	Chediak-Higashi syndrome (OMIM)
MOB4	dog	ENSG00000115540	May play a role in membrane trafficking, specifically in membrane budding reactions.	N/A	N/A
MYLK3	cat	ENSG00000140795	Kinase that phosphorylates MYL2 in vitro. Promotes sarcomere formation in cardiomyocytes and increases cardiomyocyte contractility (By similarity).	Inflammation mediated by chemokine and cytokine signaling pathway, Cytoskeletal regulation by Rho GTPase (PANTHER)	N/A

NCOA6	dog	ENSG00000198646	Nuclear receptor coactivator that directly binds nuclear receptors and stimulates the transcriptional activities in a hormone-dependent fashion. Coactivates expression in an agonist- and AF2-dependent manner. Involved in the coactivation of different nuclear receptors, such as for steroids (GR and ERs), retinoids (RARs and RXRs), thyroid hormone (TRs), vitamin D3 (VDR) and prostanooids (PPARs). Probably functions as a general coactivator, rather than just a nuclear receptor coactivator. May also be involved in the coactivation of the NF-kappa-B pathway. May coactivate expression via a remodeling of chromatin and its interaction with histone acetyltransferase proteins. Protein kinase that seems to act exclusively upon Ithreonine residues (By similarity). Required for normal entry into proliferative arrest after a limited number of cell divisions, also called replicative senescence. Required for normal cell cycle arrest in response to double-stranded DNA damage.	Circadian Clock, Developmental Biology, Gene Expression, Metabolism, Organelle biogenesis and maintenance (Reactome)	hypertensive retinopathy; malignant melanoma, somatic; breast cancer (MalaCards)
NEK4	cat	ENSG00000114904		N/A	N/A
NTSDC2	horse	ENSG00000168268		N/A	N/A
NTM	horse	ENSG00000182667	Neural cell adhesion molecule.	cellular process, developmental process (PANTHER)	childhood pilocytic astrocytoma; juvenile pilocytic astrocytoma (MalaCards)
PLACL8L1	cat, cattle	ENSG00000173261		N/A	N/A
PPAP2A	cat	ENSG000000067113	Broad-specificity phosphohydrolase that dephosphorylates exogenous bioactive glycerolipids and sphingolipids. Catalyzes the conversion of phosphatidic acid (PA) to diacylglycerol (DG). Pivotal regulator of lysophosphatidic acid (LPA) signaling in the cardiovascular system. Major enzyme responsible of dephosphorylating LPA in platelets, which terminates signaling actions of LPA. May control circulating, and possibly also regulate localized, LPA levels resulting from platelet activation. It has little activity towards ceramide-1-phosphate (C-1-P) and sphingosine-1-phosphate (S-1-P). The relative catalytic efficiency is LPA > PA > S-1-P > C-1-P. It's down-regulation may contribute to the development of colon adenocarcinoma.	Metabolism (Reactome)	N/A
PPAPDC1B	cat	ENSG00000147535	Displays magnesium-independent phosphatidate phosphatase activity in vitro. Catalyzes the conversion of phosphatidic acid to diacylglycerol. May be a metastatic suppressor for hepatocellular carcinoma.	Immune System (Reactome)	hepatocellular carcinoma (MalaCards)
PRR11	cat	ENSG000000068489	Plays a critical role in cell cycle progression. Plays a role in cell-cell adhesion through heterophilic trans-interactions with necitin-like proteins or neclins, such as trans-interaction with NECTIN2 at Sertoli-spermatid junctions. Trans-interaction with PVR induces activation of CDC42 and RAC small G proteins through common signaling molecules such as SRC and RAP1. Also involved in the formation of cell-cell junctions, including adherens junctions and synapses. Induces endocytosis-mediated down-regulation of PVR from the cell surface, resulting in reduction of cell movement and proliferation. Plays a role in the morphology of the ciliary body.	N/A	N/A
PVRL3	cattle	ENSG00000177707		Cell adhesion molecules, Adherens junction (KEGG)	N/A
RFTN2	dog	ENSG00000162944		N/A	N/A
RNPC3	cat, dog	ENSG00000185946	Participates in pre-mRNA U12-dependent splicing, performed by the minor spliceosome which removes U12-type introns. U12-type introns comprises less than 1% of all non-coding sequences. Binds to the 3'-stem-loop of m7G-capped U12 snRNA.	Gene Expression (Reactome)	growth hormone deficiency, parainfluenza virus type 3, idiopathic inflammatory myopathy, rheumatic disease (MalaCards)
SF3B1	dog	ENSG00000115524	Subunit of the splicing factor SF3B required for 'A' complex assembly formed by the stable binding of U2 snRNP to the branchpoint sequence (BPS) in pre-mRNA. Sequence independent binding of SF3A/SF3B complex upstream of the branch site is essential, it may anchor U2 snRNP to the pre-mRNA. May also be involved in the assembly of the 'E' complex. Belongs also to the minor U12-dependent spliceosome, which is involved in the splicing of rare class of nuclear pre-mRNA intron.	Spliceosome (KEGG)	Myelodysplastic syndrome, somatic (OMIM)
SKA2	dog	ENSG00000182628	Component of the SKA1 complex, a microtubule-binding subcomplex of the outer kinetochore that is essential for proper chromosome segregation. Required for timely anaphase onset during mitosis, when chromosomes undergo bipolar attachment on spindle microtubules leading to silencing of the spindle checkpoint. The SKA1 complex is a direct component of the kinetochore-microtubule interface and directly associates with microtubules as oligomeric assemblies. The complex facilitates the progressive movement of microsphears along a microtubule in a depolymerization-coupled manner. In the complex, it is required for SKA1 localization. Affinity for microtubules is synergistically enhanced in the presence of the ndc-80 complex and may allow the ndc-80 complex to track depolymerizing microtubules.	Cell Cycle, Signal Transduction (Reactome)	Post-traumatic stress disorder (MalaCards)
SNRPD1	cattle	ENSG00000167088	Core component of the spliceosomal U1, U2, U4 and U5 small nuclear ribonucleoproteins (snRNPs), the building blocks of the spliceosome. Thereby, plays an important role in the splicing of cellular pre-mRNAs. Most spliceosomal snRNPs contain a common set of Sm proteins SNRPE, SNRPF, SNRPG, SNRPH, SNRPI, SNRPD2, SNRPD3, SNRPE, SNRPF and SNRPG that assemble in a heptameric protein ring on the Sm site of the small nuclear RNA to form the core snRNP. May act as a charged protein scaffold to promote snRNP assembly or strengthen snRNP-snRNP interactions through nonspecific electrostatic contacts with RNA. Acts as a scavenger receptor for acetylated low density lipoprotein. Binds to both Gram-positive and Gram-negative bacteria and may play a role in defense against bacterial infection. When inhibited in endothelial tube formation assays, there is a marked decrease in cell-cell interactions, suggesting a role in angiogenesis. Involved in the delivery of newly synthesized CHD1/S1-CLP from the biosynthetic compartment to the endosomal/lysosomal system.	Spliceosome, Systemic lupus erythematosus (KEGG)	systemic lupus erythematosus; lupus erythematosus (MalaCards)
STAB1	horse	ENSG000000010327	May play a role in vesicle trafficking (By similarity). Binds phosphatidylinositol 3,4,5-trisphosphate. Acts as a RAS27A effector protein and may play a role in cytotoxic granule exocytosis in lymphocytes (By similarity).	Vesicle-mediated transport (Reactome)	histiocytosis; rosai-dorfman disease (MalaCards)
SYTL1	cat	ENSG00000142765		Vesicle-mediated transport (Reactome)	N/A
TAS2R16	cattle	ENSG00000128519	Gustducin-coupled receptor implicated in the perception of bitter compounds in the oral cavity and the gastrointestinal tract. Signals through PLCB2 and the calcium-regulated cation channel TRPM5.	Taste transduction (KEGG)	Alcohol dependence, susceptibility to (OMIM)
TEX14	cat	ENSG00000121101	Required both for the formation of intercellular bridges during meiosis and for kinetochore-microtubule attachment during mitosis. Intercellular bridges are evolutionarily conserved structures that connect differentiating germ cells and are required for spermatogenesis and male fertility. Acts by promoting the conversion of midbodies into intercellular bridges via its interaction with CEP55; interaction with CEP55 inhibits the interaction between CEP55 and POC5GIP1/ALX and TSG101, blocking cell abscission and leading to transform midbodies into intercellular bridges. Also plays a role during mitosis; recruited to kinetochores by PLK1 during early mitosis and regulates the maturation of the outer kinetochores and microtubule attachment. Has no protein kinase activity in vitro (By similarity).	N/A	N/A
TP53BP1	cat	ENSG000000067369	Plays a key role in the response to DNA damage. May have a role in checkpoint signaling during mitosis. Enhances TP53-mediated transcriptional activation.	NOD-like receptor signaling pathway (KEGG)	riddle syndrome; ataxia-telangiectasia; gastric cardia adenocarcinoma; fanconi anemia, complementation group a (MalaCards)
ZMYND10	cat	ENSG000000004838	Required for motile ciliary function. Probably involved in axonemal assembly of inner and outer dynein arms (IDA and ODA, respectively) for proper axoneme building for cilia motility. May act by indirectly regulating transcription of dynein proteins.	N/A	ciliary dyskinesia, primary, 22 (MalaCards)
ZNF521	cattle	ENSG00000198795	Transcription factor that can both act as an activator or a repressor depending on the context. Involved in BMP signaling and in the regulation of the immature compartment of the hematopoietic system. Associates with SMADs in response to BMP2 leading to activate transcription of BMP target genes. Acts as a transcriptional repressor via its interaction with EBF1, a transcription factor involved specification of B-cell lineage; this interaction preventing EBF1 to bind DNA and activate target genes.	N/A	N/A

Table S3: Overlapping genes between AMH and 2 or more domesticated animals

GENE NAME	SPECIES OVERLAP	ENSEMBL ID	GENE FUNCTION (UniProt)	PATHWAY ANNOTATION	RELATED DISORDERS
BRAF	cat, horse, human	ENSG00000157764	Protein kinase involved in the transduction of mitogenic signals from the cell membrane to the nucleus. May play a role in the postsynaptic responses of hippocampal neuron. Phosphorylates MAP2K1, and thereby contributes to the MAP kinase signal transduction pathway.	EGFR tyrosine kinase inhibitor resistance, Endocrine resistance, MAPK signaling pathway, ErbB signaling pathway, Rap1 signaling pathway, cAMP signaling pathway, Chemokine signaling pathway, FcγO signaling pathway, mTOR signaling pathway, Vascular smooth muscle contraction, Dorsal-ventral axis formation, Focal adhesion, Natural killer cell mediated cytotoxicity, Long-term potentiation, Neurotrophin signaling pathway, Serotonergic synapse, Long-term depression, Regulation of actin cytoskeleton, Insulin signaling pathway, Progesterone-mediated oocyte maturation, Alcoholism, Hepatitis C, Pathways in cancer, Proteoglycans in cancer, Colorectal cancer, Renal cell carcinoma, Pancreatic cancer, Endometrial cancer, Glioma, Prostate cancer, Thyroid cancer, Melanoma, Bladder cancer, Chronic myeloid leukemia, Acute myeloid leukemia, Non-small cell lung cancer, Breast cancer (KEGG); VEGF signaling pathway, T cell activation, Interleukin signaling pathway, Inflammation mediated by chemokine and cytokine signaling pathway, Integrin signalling pathway, EGF receptor signaling pathway, Ras Pathway, Angiogenesis, CCKR signaling map, PDGF signaling pathway, B cell activation (PANTHER)	Thyroid cancer; Malignant melanoma; Noonan syndrome and related disorders; Langerhans cell histiocytosis (KEGG); Adenocarcinoma of lung, somatic; Cardiofaciocutaneous syndrome; Colorectal cancer, somatic; LEOPARD syndrome; Melanoma, malignant, somatic; Non-small cell lung cancer, somatic; Noonan syndrome (OMIM)
FAM172A	cattle, dog, human	ENSG00000113391	N/A Receptor for glutamate that functions as ligand-gated ion channel in the central nervous system and plays an important role in excitatory synaptic transmission. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. The postsynaptic actions of Glu are mediated by a variety of receptors that are named according to their selective agonists. This receptor binds domoate > kainate >> L-glutamate = quisqualate >> AMPA = NMDA.	N/A	N/A
GRIK3	dog, cattle, human	ENSG00000163873	N/A Receptor for glutamate that functions as ligand-gated ion channel in the central nervous system and plays an important role in excitatory synaptic transmission. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. The postsynaptic actions of Glu are mediated by a variety of receptors that are named according to their selective agonists. This receptor binds domoate > kainate >> L-glutamate = quisqualate >> AMPA = NMDA.	Glutamatergic synapse, Neuroactive ligand-receptor interaction (KEGG); Huntington disease (PANTHER)	schizophrenia; alcohol dependence (MalaCards)
PLAC8L1	cat, cattle, human	ENSG00000173261	N/A Participates in pre-mRNA U12-dependent splicing, performed by the minor spliceosome which removes U12-type introns. U12-type introns comprises less than 1% of all non-coding sequences. Binds to the 3'-stem-loop of m7G-capped U12 snRNA.	N/A Gene Expression (Reactome)	N/A growth hormone deficiency, parainfluenza virus type 3, idiopathic inflammatory myopathy, rheumatic disease (MalaCards)
RNPC3	cat, dog, human	ENSG00000185946	N/A Participates in pre-mRNA U12-dependent splicing, performed by the minor spliceosome which removes U12-type introns. U12-type introns comprises less than 1% of all non-coding sequences. Binds to the 3'-stem-loop of m7G-capped U12 snRNA.	N/A	N/A

MFRP	NCOA6	dog	TRPC7	RAET1L	TMEM151A
MFSD11	NDUFB1	cattle	TSC22D1	RAF1	TMEM50A
MGAT4A	NEK1	cat	TSC22D4	RASSF6	TMEM87A
MB1	NEK4	cat	TSGA10	RBMS3	TMPRSS11A
MISA1	NFAH1	cat	TTC19	RC3TB2P1	TRAP
MKL1	NFKBIZ	cat	UBE2D2	REL	TRIM42
MKLN1	NID2	horse	UGDH	RG53	TSC2
MLH1	NINJ1	horse	UQCRRHP4	RHBDD3	TSNAXIP1
MMP9	NIP2L	dog	USP32	RHOA	TTC39CP1
MOB4	NIPBL	horse	USP4	RIMS2	TTC8
MPND	NKAN2	dog	USP8P1	RNF213	TT2
MRFLA9	NOCT	dog	VAV3	RNF6	TUSC2
MRPS5	NOL4	dog	VN1R66P	RPL12P41	U3
MTMR4	NOLC1	cat	VN1R67P	RPL27AP	UBE2D3P3
MTRNR2L7	NOSTRIN	cat	VRBP	RPL32	UBL5P4
MUSTN1	NOTCH2	cat	VPS41	RPL32P1	USP24P1
MYH3	NPA33	cattle	WASF5P	RPL35P6	USP43
MYH4	NPFPR2	cat	XKR3	RPL36AP40	USP46
MYL4	NPTX1	cattle	XPNPPEP3	RP56P7	USP6P1
MYLK3	NR2F2	cattle	XPDS	RSP02	VCAN
NADK2	NR3C2	horse	YWHAZP4	RTTN	VDAC1P5
NAT6	NRF1	dog	ZCWPW1	RWDD4P2	VPS13B
NAT8	NRG2	dog, cat, cattle	ZNF514	SBF2	VPS26BP1
NAV3	NRG4	cattle	ZNF568	SCARA3	VPS39
NCOA5	NRXN1	cattle	ZNF670	SCN10A	VPS8
NCOA6	NT5DC2	horse	ZNF675	SCUBE2	VWF
NDUFA11	NTAN1	dog	ZNF695	SDHAP3	WASF5P
NEAT1	NTM	horse	ZNF720	SKK1	WDRCP
NEB	NUD1T15	cat	ZNF725P	SEC11A	WR95P
NEDD1	NUMB	horse	ZNF815P	SEC63	WWP2
NEK4	NUP133	horse	ZNF92	SEMA5B	YIF1A
NETO2	NUP54	dog	ZPBP	SENP7	ZDHHC20P2
NEXN	NXPE3	dog	ZSWIM7	SERPINA3	ZDHHC24
NFG3	OLIG1	cattle		SH2D4A	ZMYM1
NISCH	OMA1	dog		SLC17A6	ZMYND10
NLK	OPML	horse		SLC25A1P2	ZNF394
NLRX1	OPTC	cat		SLC25A26	ZNF48
NMUR2	OR10K1	cat		SLC38A8	ZNF493
NOTO	OR13C8	cattle		SLC6A16	ZNF675
NPRL2	OR2B11	cat		SLC9C2	ZNF682
NR0B2	OR4D6	cat		SLFN12	ZNF747
NR2F1	OR51A7	cattle		SMARCD2	ZNF768
NT5DC2	OR8A4	dog		SMC3	ZNF771
NTM	ORF olfactory receptor family cluster (Chromo	cattle		SNRPC	ZNF789
NTRK2	ORF olfactory receptor family cluster (Chromo	cattle		SOCS2P2	
NUDC	ORF olfactory receptor family cluster (Chromo	cattle		SOQ2	
NUFIP1	ORF olfactory receptor family cluster (Chromo	cattle		SORBS1	
NUP37	OTOF	cat		SORBS2	
NWD1	PAFAH2	cat		SP140	
NWD2	PARP12	dog		SPATS2L	
NXPH1	PARVG	cat		SPDL1	
NYAP2	PCDH16	cattle		SPECC1L	
ORA5D	PCDH41	cat		SPINT3	
OTUD4	PCDH84	cat		SPOCK3	
OTX1	PCSK5	horse		SPTLC3	
PACSN1	PDE4D	dog		ST6GAL2	
PAIP2	PDE4DIP	horse		ST6GALNAC3	
PARP3	PDE5A	horse		STAB2	
PARPBP	PDE7B	dog		STAT14	
PATL2	PDLT	dog		STK31	
PBRM1	PDHR1	horse		STXBPL5L	
PC	PDXDC1	dog		TBC1D22A	
PCBP4	PEX7	cattle		TBC1D27	
PCCB	PHF2	horse		TBRG4	
PCCDH17	PHF20	horse		TDP1	
PCDH9	PHLDB3	cat		TDKXH	
PCGF6	PIK3C3	horse		TF	
PCNX	PIR1M1	cat		TF6	
PDCD4	PLA2	cat		TMEM117	
PDE4B	PKD1L1	dog		TMEM200B	
PDI3A	PLA2G2E	cat		TMEM40	
PDZD2	PLA2G3	cat		TMEM5	
PDZD3	PLAC1	cat		TMPRSS2	
PEL1	PLAC8L1	cat, cattle		TMPRSS3	
PEX13	PLAG1	cattle		TNFRSF13B	
PHACTR1	PLEC1	dog		TNFRSF1B	
PHF2	PLEKHH1	cat		TNKS1BP1	
PHKB	PLEKHM3	dog		TNP01	
PIGV	PLIN3	cat		TP53BP1	
PIK3CG	PLXNA4	horse		TPRXL	
PLA2G16	PML	cat		TPX2	
PLA2G4D	Pol	cattle		TRAT1	
PLA2G4E	POLI	dog		TRIM25	
PLA2GDF	POLR1E	dog		TRIM34	
PLAC8L1	POP1	horse		TRIM6	
PLXDC2	PPAP2A	cat		TSG101	
PMCH	PPAPDC1B	cat		TTCC23L	
POC1A	PPFIBP1	cat		TTN	
PODXL	PPM1D	horse		TUBA8	
POMGN1	PPP1R13B	cat		TUBB4BP5	
POTEC	PPP2CA	horse		U2SURP	
POU2F2	PRICKLE4	cat		US	
POU3F1	PRKAG1	cat		UBASH3A	
POU5F2	PRKCZ	horse		UTP20	
PPAP2A	PRKG2	cat		VN1R20P	
PPAPDC1A	PRMT3	horse		VN1R31P	
PPAPDC1B	PRR11	cat		VN1R32P	
PPIL4	PRX	cat		VN1R33P	
PPIP5K1	PSMB7	horse		VN2R9P	
PPM1E	PSPH	cat		VVASA	
PPM1M	PSTK	cat		WDF4	
PPP2R1B	PTPN4	horse		WDR47	
PRADC1	PTPRR	cat		WDR62	
PRDM10	PTPRS	cat		WDR87	
PRDM2	PUSL1	cat		XKR3	
PRKAR2B	PVRL3	cattle		YTHDF2	
PRKDC	Q2ABD2	dog		ZDHHC13	
PRKDD	RAB3GAP1	dog		ZDHHC20P2	
PRKME2	RABGAP1L	dog		ZNF280D	
PRR11	RABL3	cat		ZNF33BP1	
PSTPIP2	RALY	dog		ZNF592	
PTPN23	RANBP17	dog		ZNF605	
PTPRD	RAPH1	dog		ZNF7	
PUS10	RBM11	cat		ZNF718	
PVRL3	RBP5	cat		ZNF804B	
QSER1	RCS01	cat		ZNF850	
RAB11F1P5	REEP1	dog		ZNF861P	
RAB28	RELL1	cat		ZSWIM5	
RABAC1	RELT	dog			
RAD51C	RFTN2	dog			
RAD54L	RG9MTD3	dog			
RAD54L2	RHBDD1	cat			
RANBP1	RHPN1	horse			
RANBP3	RIMKLA	cat			
RARRES3	RNAS5E	cat			
RASA1	RNF103	dog			
RASGEF1A	RNF144B	cattle			
RASSF1	RNPC3	dog, cat			
RASSF3	ROBO1	cattle			
RB1CC1	RPL3	dog			

RBF0X2	RPL31	dog
RBL1	RRN3	dog
RBM14	RRN3P1	dog
RBM15B	RRNRP2	dog
RBM4	RSL1D1	cat
RBM4B	RTP3	cat
RBSG3	S100A12	cat
RCE1	SAE1	dog
REL	SCARB2	dog
RET	SCN9A	cat
RFT1	SCP2D1	dog
RFTM2	SCOPE1	horse
RC36	SCRIB	cat
RIF1	SDAD1	dog
RNF133	SDK2	cat
RNF148	SEC24A	cat, horse
RNF220	SEC63	horse
RNF26	SEMA3D	dog
RNF43	SEM6A	cattle
RNF44	SENP5	cat
RNPC3	SENP7	cat
ROBO2	SEPT10	cat
ROCK1	SERINC3	cat
RPL13AP6	SETBP1	dog
RPL29	SETD9	dog
RPS18P9	SETMAR	cattle
RPS6K6A1	SF3B1	dog
RBP9	SGCD	horse
RSP03	SH2D5	cat
SAMHD1	SH3GL2	dog
SCAP	SHC4	cat
SCMH1	SIAE	cat
SCYL3	SKA2	dog
SEC23IP	SKP1	horse
SEC24D	SLC22A13	cat
SEMA3F	SLC22A15	horse
SEMA3G	SLC22A18	cat
SEMA6D	SLC25A38	cat
SEPT4	SLC35D1	cattle
SERF2	SLC35F5	cat
SERINC4HYPK	SLC39A7	cat
SESN1	SLC39A8	cat
SETD1A	SLC41A2	cattle
SFA3	SLC43A1	horse
SF3B1	SLC46A1	cat
SFMBT1	SLC5A1	dog
SFXN5	SLC5A4	dog
SGMS2	SLC6A1	cattle
SGSM3	SLC6A17	dog
SH3GL1	SLC9A6	dog
SH3RF2	SLC01A2	cat
SHOC2	SMC4	dog
SHZ	SMC1	cat
SIN3B	SMG6	cat, horse
SIPA1L1	SMIM23	dog
SKA2	SMO	dog
SKIV2L2	SMYD2	dog
SKP2	SNAP29	dog
SLC12A5	SNRPD1	cattle
SLC16A1	SPATA19	horse
SLC1A1	SPATA21	cat
SLC25A17	SPATA7	cat
SLC26A3	SPERT	cat
SLC26A4	SPHKAP	cat
SLC2A5	SPINT1	cat
SLC30A2	SPTAN1	cattle
SLC35B1	SPTBNS	cat
SLC35E1	SREBF1	cat
SLC38A9	SRF72	dog
SLC4A10	SRRM2	cat
SLC4A4	STAB1	horse
SLC9A1	STARDS	cat
SLFNL1	STARDS	dog
SLIT2	STK10	dog, cattle
SLITR1	STK11P	cat
SLITR3	STS	cat
SMAD1	STX7	dog
SMAD9	STXBP6	horse
SMG8	SUN3	cat
SMIM4	SURF2	cat
SMIM7	SUSD3	horse
SMYD5	SYNM	cat
SNAI2	SYTL1	cat
SNAP23	TACK1	cattle
SNF	TAS2R1	cat
SNHG4	TAS2R16	cattle
SNRPD1	TAS2R3	cat
SORCS1	TAS2R38	dog
SORCS2	TBC1D9	dog
SORCS3	TBXAS1	dog
SPATS2L	TCTN1	horse
SPCS1	TCTN3	dog
SPG11	TEK3	dog
SPIDR	TEX14	cat
SPOP	TF	cat
SPTB2	TFPC2L1	cattle
SRSF2	TH	dog
ST7	THBS2	cat
STAB1	THEGL	dog
STAD	THUMPD1	cat
STAG1	THYN1	horse
STAMPB	TLX3	dog
STARSD	TMEM114	dog
STK3	TMEM132D	dog, cattle
STMN2	TMEM159	dog
STRC	TMEM242	dog
STX1A	TMEM59L	cat
STX1B	TMEM71	cat
SUP74H1	TNRSF9	cattle
SYNP02	TNKS2	dog
SYT1	TOE1	cat
SYT6	TP53BP1	cat
SYTL1	TRAPPC8	horse
SYVN1	TRBV25OR9-2	dog
TAC4	TRIM16	dog
TAF5	TRIM59	dog
TANC2	TRIO	horse
TAS2R16	TRMT61A	dog
TBC1D23	TRPV6	cat
TBX1	TRY1	dog
TDRD3	TRY2	dog
TDRD7	TRY3	dog
TEX14	TSTD2	dog
TEX264	TTC38A	dog
TFA2D	TUBGCP5	dog
TGM4	TVP23B	dog
TGM5	TVP23C	dog
TGM7	TXN2	cat
THSD7B	TXNRD2	cat
THTPA	TYK2	cat

TK2	U2	dog
TKT	UBE2B	horse
TLE3	UBXN10	cat
TLR9	ULBP3	cattle
TMTSF2	UMOD	dog
TMEM110	UNC93A	cattle
TMEM115	URB2	horse
TMEM123	USP45	cat
TMEM17	UVRAG	cat
TMEM222	V1R	dog
TMEM235	VDAC1	horse
TMEM262	VEZT	dog, cat
TMEM38A	VPS26B	horse
TMEM42	VRK1	horse
TMEM62	VWC2	dog
TMEM67A	VWDE	cattle
TMOD1	WASF3	horse
TNFRSF21	WDR17	cat
TNRC1	WDR62	cat
TNRC9B	WDR90	cat
TNS1	WFDC9	cat
TP53BP1	WIPF2	cat
TP53INP2	WIPF3	cat
TPD52	WNK2	horse
TPRKB	WWC1	cat
TRANK1	XCR1	cat
TRIM37	XPBP	dog
TRIM43	XP-C	cat
TRIM69	XPO6	cattle
TRIM71	YSK4	dog
TRMT2A	YWHAH	dog
TRNP1	ZC3H3	horse
TTBK2	ZFAT	cat
TTC6	ZFYVE19	cat
TUBGCP4	ZMYND10	cat
TUSC2	ZNF236	dog
TWF2	ZNF286A	dog
TYW5	ZNF286B	dog
U6	ZNF436	cat
U7	ZNF492	dog
UBE2V2	ZNF516	dog
UBR1	ZNF521	cattle
UGGT2	ZNF555	cat
UGP2	ZNF622	cat
UGT8	ZNF679	dog
UNC50	ZNF780B	cat
UOCRH	ZP2	dog
UOCRHL	ZPBP	dog
USP33	ZZEF1	cat
USP34		
USP54		
UTP11		
VAFB		
VMAC		
VOPP1		
VPS39		
VPS51		
VPS54		
WASF2		
WISSCR22		
WDPCP		
WDR59		
WDR76		
WDR82		
WDT01		
WHSC1L1		
WIZ		
XP01		
ZBBX		
ZBTB20		
ZBTB34		
ZDHHC18		
ZDHHC8		
ZEB2		
ZFHx4		
ZFL1		
ZIC4		
ZMYND10		
ZNF106		
ZNF197		
ZNF2		
ZNF205		
ZNF213		
ZNF248		
ZNF25		
ZNF33A		
ZNF33B		
ZNF35		
ZNF37A		
ZNF407		
ZNF501		
ZNF502		
ZNF514		
ZNF521		
ZNF574		
ZNF638		
ZNF852		
ZNHIT2		
ZNRF1		
ZSCAN29		

Table S4b: Contrasts between domesticates, between AMH and each domesticate, and between great apes and domesticates (for further statistical analysis, see Table S4c)

	dog = 229	cat = 291	cattle = 78	horse = 109	AMH = 742
dog = 229		$v = 5, p = 0.2575646$	$v = 5, p = 0.002250835$	$v = 0$	$v = 15, p = 0.02334542$
cat = 291	$v = 5, p = 0.2575646$		$v = 3, p = 0.1111013$	$v = 4, p = 0.0805177$	$v = 15, p = 0.1454225$
cattle = 78	$v = 5, p = 0.002250835$	$v = 3, p = 0.1111013$		$v = 0$	$v = 9, p = 0.002784603$
horse = 109	$v = 0$	$v = 4, p = 0.0805177$	$v = 0$		$v = 7, p = 0.1219762$

$p < 0.05$
 $p < 0.01$

Intersections between domesticates				Intersections between total AMH and each domesticate				Intersections between great apes and domesticates			
DOG-CAT [5]	DOG-CATTLE [5]**	CAT-CATTLE [3]	CAT-HORSE [4]	AMH-DOG [15]*	AMH-CAT [15]	AMH-CATTLE [9]**	AMG-HORSE [7]	DOM-CHIMP [16]	DOM-ORANGO [20]	DOM-GORILLA [12]	
ATXN7L1	FAM172A	ADAMT513	BRAP	COA5	BRAP	ERBB4	AMBRA1	ALK	ACOX3	CUX2	
CLECSA	GRK3	NRG2	DCC	COL11A1	GRIK1	FAM172A	BRAP	AMBRA1	ADAMDEC1	DNAH9	
NRG2	NRG2	PLAC8L1	SEC24A	COQ10B	HS03B7	GRIK3	CACNA1D	C3orf62	ADAMTSL3	FER	
RNPC3	STK10		SMG6	FAM172A	ITGA9	LRP1B	DLGAP1	EHBP1L1	ALK	IMMP2L	
VEZT	TMEM132D			GGT7	MYLK3	PLAC8L1	NTSDC2	FAF1	BARO1	MRPL11	
				GRK3	NEK4	PVRL3	NTM	HADH	DNAJB9	OPCML	
				HSPD1	PLAC8L1	SNRPD1	STAB1	INPP4B	FAM172A	cattle_dog	
				HSPE1	PPAP2A	TAS2R16		LCAT	ISG15	PRKAG1	
				LYST	PPAPDC1B	ZNF521		LHFPL3	KDM3A	RSL1D1	
				MCS4	PRR11			MGAM	KYNU	SCNA9A	
				NCOA6	RNPC3			NIPBL	NRXN1	SGCD	
				RFTN2	SYTL1			NXPE3	PITRM1	ZMYND10	
				RNPC3	TEX14			RABGAP1L	PLCE1		
				SFB1	TP53BP1			STABP6	POLI		
				SKA2	ZMYND10			TRBV25OR3-2	RABGAP1L		
								ZBPB	SEC63		
									SENP7		
									TF		
									TP53BP1		
									WDR62		

* $p < 0.05$
 ** $p < 0.01$

Table S4c: Results of the hypergeometric intersection tests

		Domesticates				Total domesticates = 691
		Dog = 229	Cat = 291	Cattle = 78	Horse = 109	
AMH	Peyrégne et al. 2016 = 375	-	-	-	-	$v = 18; p = 0.1199135$
	Racimo 2016 = 326	-	-	-	-	$v = 16; p = 0.1191519$
	Prüfer et al. 2014 = 108	-	-	-	-	$v = 9; p = 0.01456058 *$
	Prüfer et al. 2014 + Racimo 2016 = 419	-	-	-	-	$v = 24; p = 0.01478488 *$
	Total AMH (Peyrégne et al. 2016, Racimo 2016, Prüfer et al. 2014) = 742	$v = 15; p = 0.02934342 *$	$v = 15; p = 0.1454225$	$v = 9; p = 0.002784603 **$	$v = 7; p = 0.1219762$	$v = 41; p = 0.003407041 **$
Great apes (Cagan et al. 2016)	Chimpanzee (<i>Pan t. troglodytes</i>) = 415	$v = 6; p = 0.3614671$	$v = 5; p = 0.7449666$	$v = 0$	$v = 5; p = 0.08336393$	$v = 16; p = 0.4008807$
	Orango (<i>Pongo abelii</i>) = 500	$v = 5; p = 0.7023469$	$v = 12; p = 0.07285972$	$v = 2; p = 0.598045$	$v = 2; p = 0.7728682$	$v = 20; p = 0.3207204$
	Gorilla (<i>G. g. gorilla</i>) = 426	$v = 2; p = 0.9619034$	$v = 7; p = 0.4519085$	$v = 0$	$v = 3; p = 0.4267467$	$v = 12; p = 0.8291688$