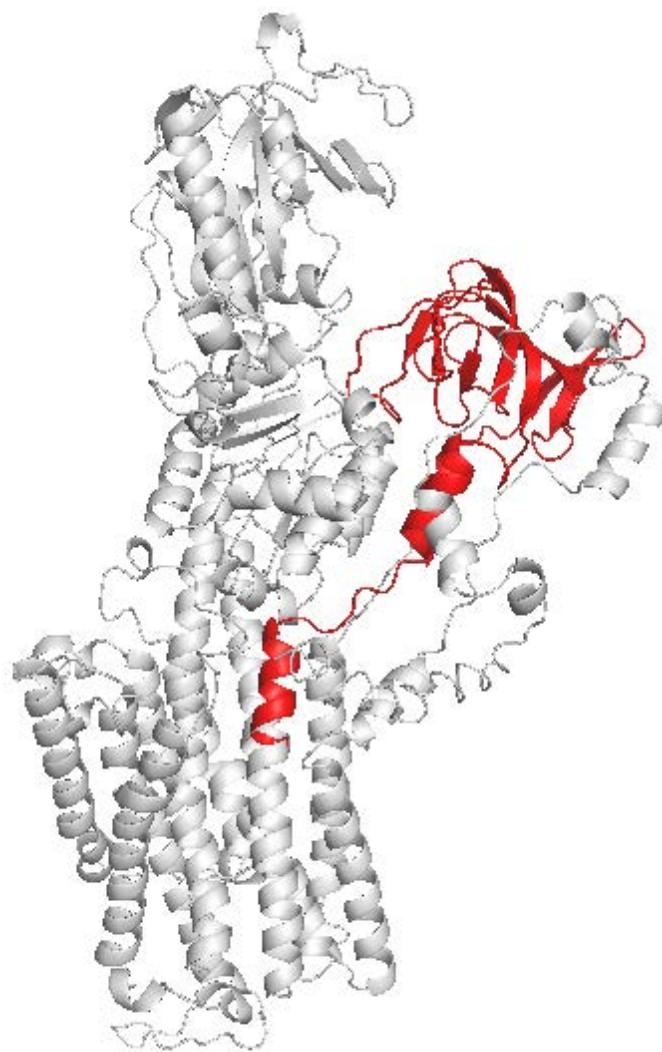


# The Non-Gastric H<sup>+</sup>/K<sup>+</sup> ATPase (ATP12A) Is Expressed in Mammalian Spermatozoa

## Supplementary Material



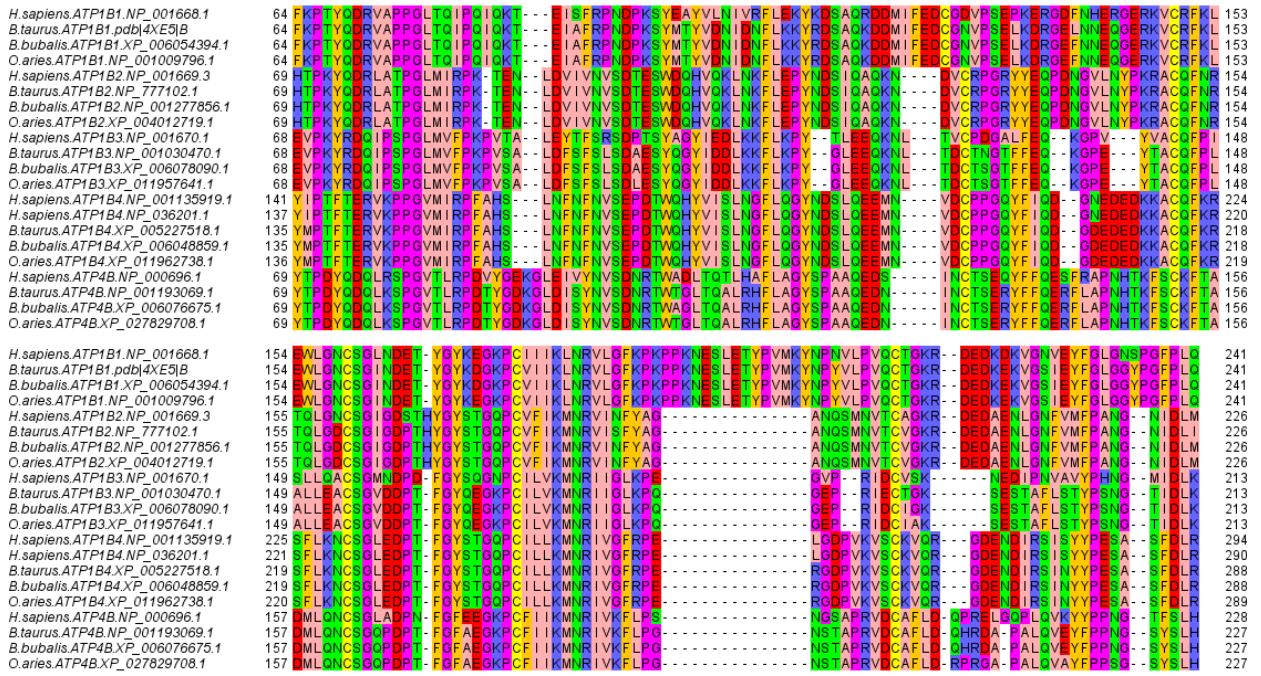
**Figure S1.** The 3D comparative model of bovine ATP12A is reported in white cartoon representation. Red cartoon indicates the localization of residues 171-306, which form the target region of the employed A62134 EpiGentek antibody.

Homo.sapiens.ATP12A.i1.NP_001667	171 NIMSSFKNM I PQQALV R OSEKKT T SEGLVVGDI VEVKGCD I PAD IRVLSSCGCRVDNSSL T CESEP PQRS	243
Homo.sapiens.ATP12A.i2.NP_001172014	171 NIMSSFKNM I PQQALV R OSEKKT T SEGLVVGDI VEVKGGDQ I PAD IRVLSSCGCRVDNSSL T CESEP PQRS	243
Bos.taurus.ATP12AXP_002691916	169 NIMSSFRKMI PQQALV R OSEKKT T PADOLVVGDI VEIKGGDRVPADRL STOGCKVDNSSL T CESEP PQRS	241
Bubalus.bubalis.ATP12AXP_006046156	170 NIMSSFRKMI PQQALV R OSEKKT T PADOLVVGDI VEIKGGDRVPADRL STOGCKVDNSSL T CESEP PQRS	242
Ovis.aries.ATP12AXP_004012362	191 NIMSSFRKMI PQQALV R OSEKKT T PADOLVVGDI VEIKGGDRVPADRL STOGCKVDNSSL T CESEP PQRS	242
Homo.sapiens.ATP4ANP_000695	167 NIIASFKNLVLPQOATV R RQCDKFQ I NADOLVVGDL VEIKGGDRVPADRL I LAAGCKVDNSSL T CESEP PQRS	239
Bos.taurus.ATP4ANP_001137561	166 NIIASFKNLVLPQOATV R RQCDKFQ I NADOLVVGDL VEMKGGDRVPADRL I LAAGCKVDNSSL T CESEP PQRS	238
Bubalus.bubalis.ATP4AXP_006068866	166 NIIASFKNLVLPQOATV R RQCDKFQ I NADOLVVGDL VEMKGGDRVPADRL I LAAGCKVDNSSL T CESEP PQRS	238
O.aries.ATP4AXP_027833250.1	155 NIIASFKNLVLPQOATV R RQCDKFQ I NADOLVVGDL VEMKGGDRVPADRL I LAAGCKVDNSSL T CESEP PQRS	217
Homo.sapiens.ATP1A1.NP_000692	145 KIMESFKNMVPPQOALV R RECEKMS I NAEEVVVGDLY VEKGGRIPADRL I SANGCKVDNSSL T CESEP PQRS	228
Bos.taurus.ATP1A1.NP_001070266	154 KIMESFKNMVPPQOALV R RECEKMS I NAEEVVVGDLY VEKGGRIPADRL I SANGCKVDNSSL T CESEP PQRS	226
B.bubalis.ATP1A1.XP_006058746.1	154 KIMESFKNMVPPQOALV R RECEKMS I NAEEVVVGDLY VEKGGRIPADRL I SANGCKVDNSSL T CESEP PQRS	226
Ovis.aries.ATP1A1.XP_027824133	125 KIMESFKNMVPPQOALV R RECEKMS I NAEEVVVGDLY VEKGGRIPADRL I SANGCKVDNSSL T CESEP PQRS	197
Homo.sapiens.ATP1A2.NP_000693	154 KIMDSFKNMVPPQOALV R RECEKMO I NAEEVVVGDLY VEKGGRIPADRL I SSHGCKVDNSSL T CESEP PQRS	226
Bos.taurus.ATP1A2.NP_001074993.1	154 KIMDSFKNMVPPQOALV R RECEKMO I NAEEVVVGDLY VEKGGRIPADRL I SSHGCKVDNSSL T CESEP PQRS	226
B.bubalis.ATP1A2.XP_006060545.1	154 KIMDSFKNMVPPQOALV R RECEKMO I NAEEVVVGDLY VEKGGRIPADRL I SSHGCKVDNSSL T CESEP PQRS	226
Ovis.aries.ATP1A2XP_004002710	154 KIMDSFKNMVPPQOALV R RECEKMO I NAEEVVVGDLY VEKGGRIPADRL I SSHGCKVDNSSL T CESEP PQRS	226
Homo.sapiens.ATP1A3.NP_689509	146 KIMESFKNMVPPQOALV R RECEKMO I NAEEVVVGDLY VEKGGRIPADRL I SAHCKVDNSSL T CESEP PQRS	218
Bos.taurus.ATP1A3.XP_024834678.1	160 KIMESFKNMVPPQOALV R REGEKMO I NAEEVVVGDLY VEKGGRIPADRL I SAHCKVDNSSL T CESEP PQRS	232
B.bubalis.ATP1A3.XP_006041318.1	160 KIMESFKNMVPPQOALV R REGEKMO I NAEEVVVGDLY VEKGGRIPADRL I SAHCKVDNSSL T CESEP PQRS	232
O.aries.ATP1A3.XP_027834157.1	147 KIMESFKNMVPPQOALV R REGEKMO I NAEEVVVGDLY VEKGGRIPADRL I SAHCKVDNSSL T CESEP PQRS	219
Homo.sapiens.ATP1A4.NP_653300	164 KIMESFKNMVPPQOALV R RGEEKMO I NAEVVVLGDL VEKGGRIPADRL I SANGCKVDNSSL T CESEP PQRS	236
Bos.taurus.ATP1A4.NP_001137575	165 KIMESFKNMVPPQOALV R RGEEKMO I PVFEYVVVGDL VEKGGRIPADRL I SSCGCKVDNSSL T CESEP PQRS	237
B.bubalis.ATP1A4.XP_006060547.1	165 KIMESFKNMVPPQOALV R RGEEKMO I PVFEYVVVGDL VEKGGRIPADRL I SSCGCKVDNSSL T CESEP PQRS	237
O.aries.ATP1A4.XP_014947816.1	165 KIMESFKNMVPPQOALV R RGEEKMO I PVFEYVVVGDL VEKGGRIPADRL I SSCGCKVDNSSL T CESEP PQRS	237
Homo.sapiens.ATP12A.i1.NP_001667	244 SEFTTHEPLETKNI CFYSTTCLEG --- T TV GMVI NT GDR I GH IA LASGVCGNEKTP IA IE IE E HF	306
Homo.sapiens.ATP12A.i2.NP_001172014	244 SEFTTHEPLETKNI CFYSTTCLEG --- T TV GMVI NT GDR I GH IA LASGVCGNEKTP IA IE IE E HF	306
Bos.taurus.ATP12AXP_002691916	242 CEFTHESPLETKNI AFFSTTCLEG --- TA GMVI NT GDR I GH IA LASGVCGNEKTP IA IE IE E HF	304
Bubalus.bubalis.ATP12AXP_006046156	243 CEFTHESPLETKNI AFFSTTCLEG --- TA GMVI NT GDR I GH IA LASGVCGNEKTP IA IE IE E HF	305
Ovis.aries.ATP12AXP_004012362	264 CEFTHESPLETKNI AFFSTTCLEG --- TA GMVI NT GDR I GH IA LASGVCGNEKTP IA IE IE E HF	326
Homo.sapiens.ATP4ANP_000695	240 PECTHESPLETRNI AFFSTMCLEG --- TV GLVVNT GDR I GH IA LASGVENEKTP IA IE IE E HF	302
Bos.taurus.ATP4ANP_001137561	239 PECTHESPLETRNI AFFSTMCLEG --- TA GLVVNT GDR I GH IA LASGVENEKTP IA IE IE E HF	301
Bubalus.bubalis.ATP4AXP_006068866	239 PECTHESPLETRNI AFFSTMCLEG --- TA GLVVNT GDR I GH IA LASGVENEKTP IA IE IE E HF	301
O.aries.ATP4AXP_027833250.1	218 PDCTHESPLETRNI AFFSTMCLEG PLGPPAGT TA GLVVNT GDR I GH IA LASGVENEKTP IA IE IE E HF	289
Homo.sapiens.ATP1A1.NP_000692	229 PDFTNENPLETRNI AFFSTNCVEG --- TARG IVVY T GDR T VMGR IA LASGLEGGQT IA IE IE E HF	291
Bos.taurus.ATP1A1.NP_001070266	227 PDFTNENPLETRNI AFFSTNCVEG --- TARG IVVY T GDR T VMGR IA LASGLEGGQT IA IE IE E HF	289
B.bubalis.ATP1A1.XP_006058746.1	227 PDFTNENPLETRNI AFFSTNCVEG --- TARG IVVY T GDR T VMGR IA LASGLEGGQT IA IE IE E HF	289
Ovis.aries.ATP1A1.XP_027824133	198 PDFTNENPLETRNI AFFSTNCVEG --- TARG IVVY T GDR T VMGR IA LASGLEGGQT IA IE IE E HF	260
Homo.sapiens.ATP1A2.NP_000693	227 PEFTTHEPLETRNI CFFSTNCVEG --- TARG IVVIA T GDR T VMGR IA LASGLEVGRTP IA IE IE E HF	289
Bos.taurus.ATP1A2.NP_001074993.1	227 PEFTTHEPLETRNI CFFSTNCVEG --- TARG IVVIA T GDR T VMGR IA LASGLEVGRTP IA IE IE E HF	289
B.bubalis.ATP1A2.XP_006060545.1	227 PEFTTHEPLETRNI CFFSTNCVEG --- TARG IVVIA T GDR T VMGR IA LASGLEVGRTP IA IE IE E HF	289
Ovis.aries.ATP1A2XP_004002710	227 PEFTTHEPLETRNI CFFSTNCVEG --- TARG IVVIA T GDR T VMGR IA LASGLEVGRTP IA IE IE E HF	289
Homo.sapiens.ATP1A3.NP_689509	219 PDOTHENPLETRNI TFFSTNCVEG --- TARGVVVA T GDR T VMGR IA LASGLEVKTP IA IE IE E HF	281
Bos.taurus.ATP1A3.XP_024834678.1	233 PDCTHDNPLETRNI TFFSTNCVEG --- TARGVVVA T GDR T VMGR IA LASGLEVKTP IA IE IE E HF	295
B.bubalis.ATP1A3.XP_006041318.1	233 PDCTHDNPLETRNI TFFSTNCVEG --- TARGVVVA T GDR T VMGR IA LASGLEVKTP IA IE IE E HF	295
O.aries.ATP1A3.XP_027834157.1	220 PDCTHDNPLETRNI TFFSTNCVEG --- TARGVVVA T GDR T VMGR IA LASGLEVKTP IA IE IE E HF	282
Homo.sapiens.ATP1A4.NP_653300	237 PDFTHEPLETRNI CFFSTNCVEG --- TARGIVIA T GDS T VMGR IA LTSGLAVGQTP IA IE IE E HF	299
Bos.taurus.ATP1A4.NP_001137575	238 TEFTNENPLETRNI CFFSTNCVEG --- SAGIVIA T GDS T VMGR IA LTSGLAVGQTP IA IE IE E HF	300
B.bubalis.ATP1A4.XP_006060547.1	238 TEFTNENPLETRNI CFFSTNCVEG --- SAGIVIA T GDS T VMGR IA LTSGLAVGQTP IA IE IE E HF	300
O.aries.ATP1A4.XP_014947816.1	238 TEFTNENPLETRNI CFFSTNCVEG --- SAGIVIA T GDS T VMGR IA LTSGLAVGQTP IA IE IE E HF	300

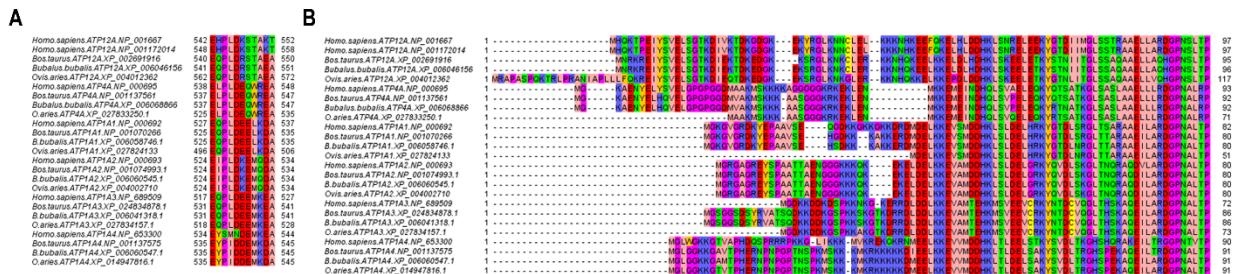
**Figure S2.** MSA of residues from the different sampled ATP1A1-4, ATP12A and ATP4A sequences, aligned with 171-306 residues from the bovine ATP12A, forming the target region of the employed A62134 EpiGentek antibody.



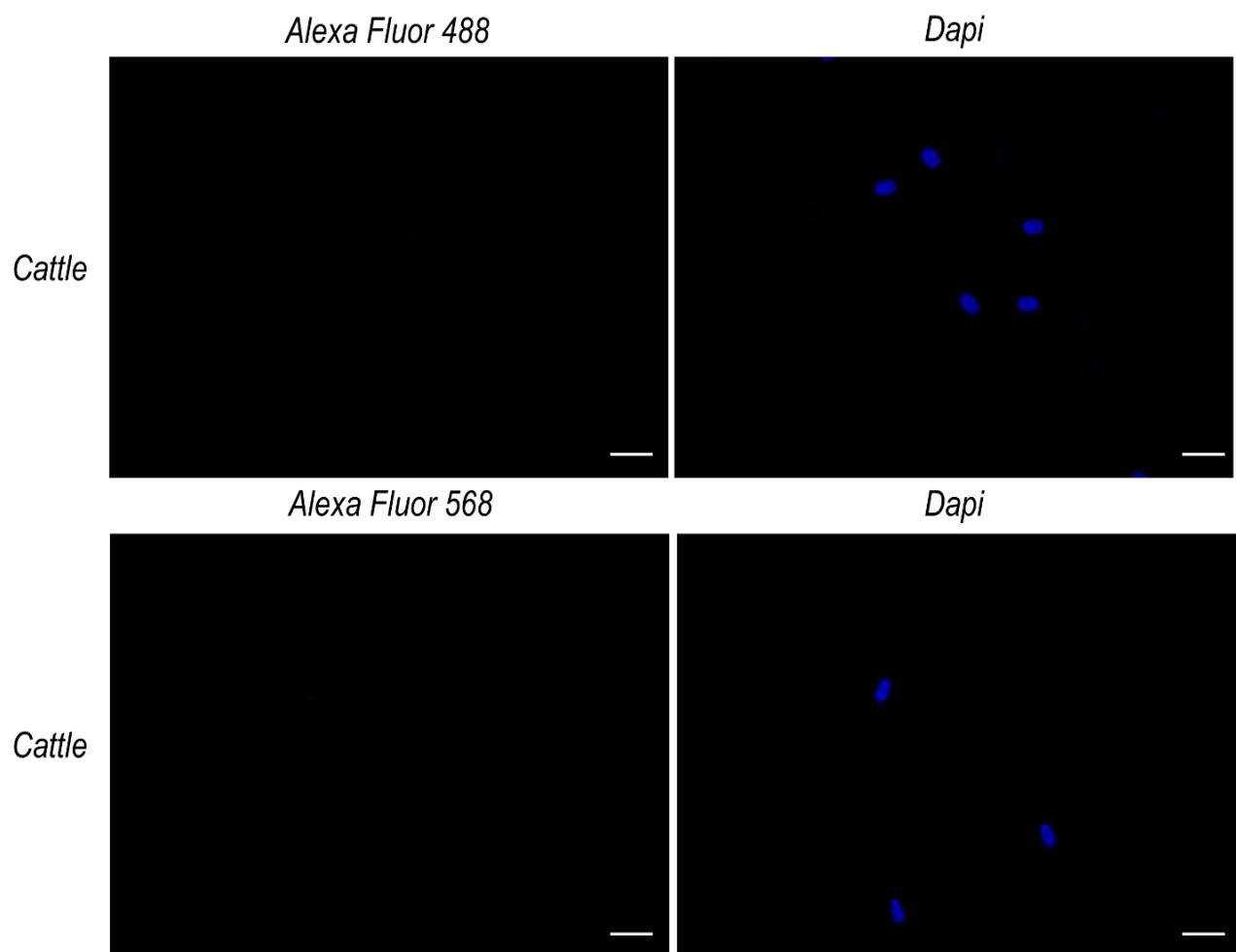
**Figure S3.** The 3D crystallized structure of bovine ATP1A1 (white cartoon representation) in complex with ATP1B1 (black cartoon representation) is reported. Red cartoon indicates the localization of residues 64-241, which form the target region of the employed CSB-PA002326LA01HU antibody.



**Figure S4.** MSA of residues from the different sampled ATP1B1-4 and ATP4B sequences, aligned with 64-241 residues from the bovine ATP1B1, forming the target region of the employed CSB-PA002326LA01HU antibody.



**Figure S5.** (A) MSA of residues from the different sampled ATP1A1-4, ATP12A and ATP4A sequences, aligned with 496-506 residues from lamb ATP1A1, forming the target region of the employed MA3-929 Invitrogen antibody. (B) MSA of residues from the different sampled ATP1A1-4, ATP12A and ATP4A sequences, aligned with 496-506 residues from the lamb ATP1A1, forming the target region of the employed MA3-929 Invitrogen antibody.



**Figure S6.** Negative controls sperms were incubated with either anti-rabbit or anti-mouse IgG secondary antibodies conjugated to Alexa Fluor 488 (upper panels) or Alexa Fluor 568 (lower panels), respectively. Nuclei were counterstained with DAPI (blue). Scale bar: 20  $\mu$ m.