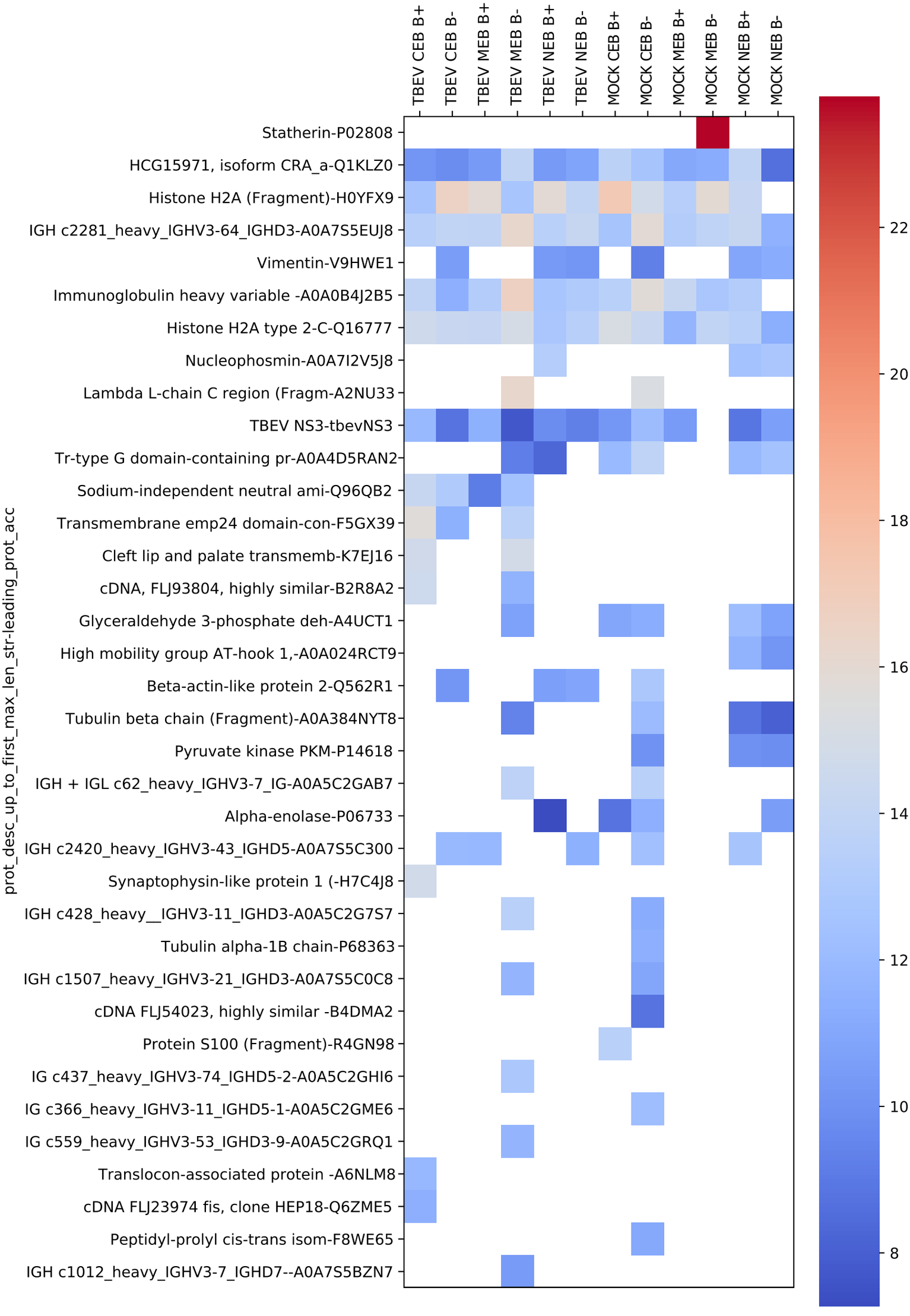
TBEV CEB B+
TBEV CEB B-
TBEV MEB B+
TBEV MEB B-
TBEV NEB B+
TBEV NEB B-
MOCK CEB B+
MOCK CEB B-
MOCK MEB B+
MOCK MEB B-
MOCK NEB B+
MOCK NEB B-
Statherin-P02808
HCG15971, isoform CRA_a-Q1KLZ0
Histone H2A (Fragment)-H0YFX9
IGH c2281_heavy_IGHV3-64_IGHD3-A0A7S5EUJ8
Vimentin-V9HWE1
Immunoglobulin heavy variable -A0A0B4J2B5
Histone H2A type 2-C-Q16777
Nucleophosmin-A0A7I2V5J8
Lambda L-chain C region (Fragm-A2NU33
TBEV NS3-tbevNS3
Tr-type G domain-containing pr-A0A4D5RAN2
Sodium-independent neutral ami-Q96QB2
Transmembrane emp24 domain-con-F5GX39
Cleft lip and palate transmemb-K7EJ16
cDNA, FLJ93804, highly similar-B2R8A2
Glyceraldehyde 3-phosphate deh-A4UCT1
High mobility group AT-hook 1,-A0A024RCT9
Beta-actin-like protein 2-Q562R1
Tubulin beta chain (Fragment)-A0A384NYT8
Pyruvate kinase PKM-P14618
IGH + IGL c62_heavy_IGHV3-7_IG-A0A5C2GAB7
Alpha-enolase-P06733
IGH c2420_heavy_IGHV3-43_IGHD5-A0A7S5C300
Synaptophysin-like protein 1 (-H7C4J8
IGH c428_heavy__IGHV3-11_IGHD3-A0A5C2G7S7
Tubulin alpha-1B chain-P68363
IGH c1507_heavy_IGHV3-21_IGHD3-A0A7S5C0C8
cDNA FLJ54023, highly similar -B4DMA2
Protein S100 (Fragment)-R4GN98
IG c437_heavy_IGHV3-74_IGHD5-2-A0A5C2GHI6
IG c366_heavy_IGHV3-11_IGHD5-1-A0A5C2GME6
IG c559_heavy_IGHV3-53_IGHD3-9-A0A5C2GRQ1
Translocon-associated protein -A6NLM8
cDNA FLJ23974 fis, clone HEP18-Q6ZME5
Peptidyl-prolyl cis-trans isom-F8WE65
IGH c1012_heavy_IGHV3-7_IGHD7--A0A7S5BZN7
prot_desc_up_to_first_max_len_str-leading_prot_acc
22
20
18
16
14
12
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8