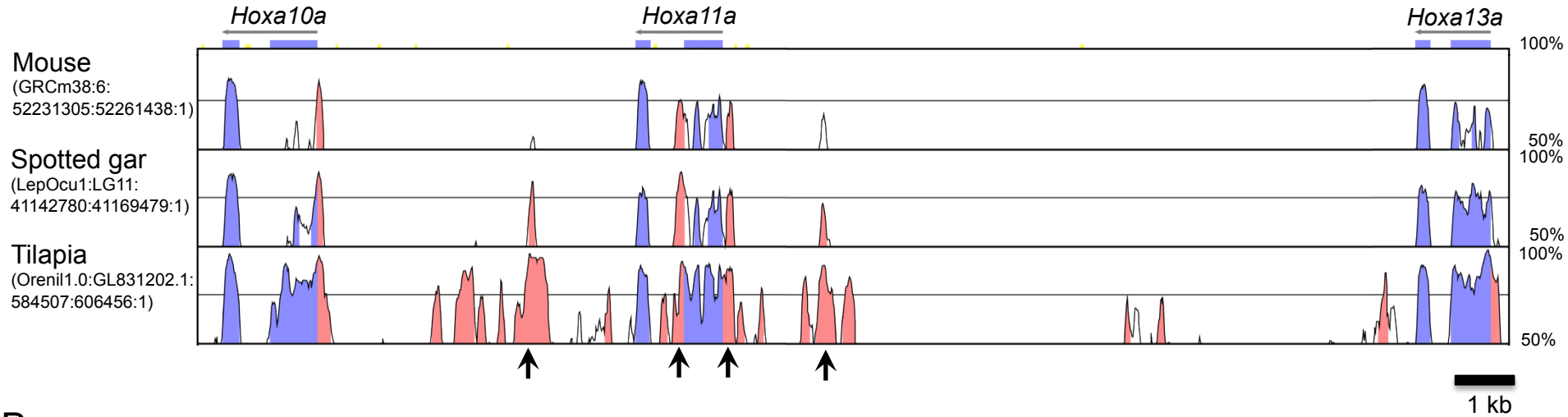


A

Tetraodon
(TETRAODON8:21:3002595:3024897:-1)



B

Length: 325
Identity: 196/325 (60.3%)
Similarity: 225/325 (69.2%)
Gaps: 56/325 (17.2%)
Score: 919.0

mmu-HOXA11	1	M M D F D E R G P C S S N M Y L P S C T Y Y V S - G P D F S S L P S F L P Q T P S S R P M T Y S Y S	49
tni-HOXA11A	1	M M D F D E R V P V G S N M Y L P S C T Y Y V S A G A D F S G L P S F - - - - S T R P M T Y S Y S	45
mmu-HOXA11	50	S N L P O V Q P V R E V T F R E Y - A I E P A T K W H P R G N L A H C Y - S A E E L V H R D C L Q A	97
tni-HOXA11A	46	S N L P O V Q P V R E V T F R D Y A A I D A S G K W S H R G N L P Q C Y P S A E D M V H R D C L S A	95
mmu-HOXA11	98	P S A A G V P G D V L A K S S - - A N V Y H H - - - - - P T P A V S S N F Y S T V G R N G V L P Q	139
tni-HOXA11A	96	Q T A - - V V G D M P G K N S P A A A A Y H H H H H H H A S A G S A A T N F Y G N V G R N G V L P Q	143
mmu-HOXA11	140	A F D Q F F E T A Y G - T P E N L A S S D Y P G D K N A E K G P Q A A A A T S A A A V A A A A T G A	188
tni-HOXA11A	144	A F D Q F Y D T A Y G N A S E N S S T N G F S G D K T P T K - - Q A S S A P E D A Q - - - - -	183
mmu-HOXA11	189	P A T S S S D G G G G G C Q E A A A A E K E R R R R P E S S S S P E S S G H T E D K A G G S G G	238
tni-HOXA11A	184	- - - - - A C R D T G A K E P R - - - - - E E T S S P E L S - - - - - S D G	206
mmu-HOXA11	239	Q R T R K K R C P Y T K Y Q I R E L E R E F F S V Y I N K E K R L Q L S R M L N L T D R Q V K I W	288
tni-HOXA11A	207	Q R T R K K R C P Y S K Y Q I R E L E R E F F S V Y I N K E K R M Q L S R M L N L T D R Q V K I W	256
mmu-HOXA11	289	F Q N R R M K E K K I N R D R L Q Y Y S A N P L L	313
tni-HOXA11A	257	F Q N R R M K E K K L N R D R L Q Y Y T S N P L L	281

DUF3528

homeodomain

1 kb