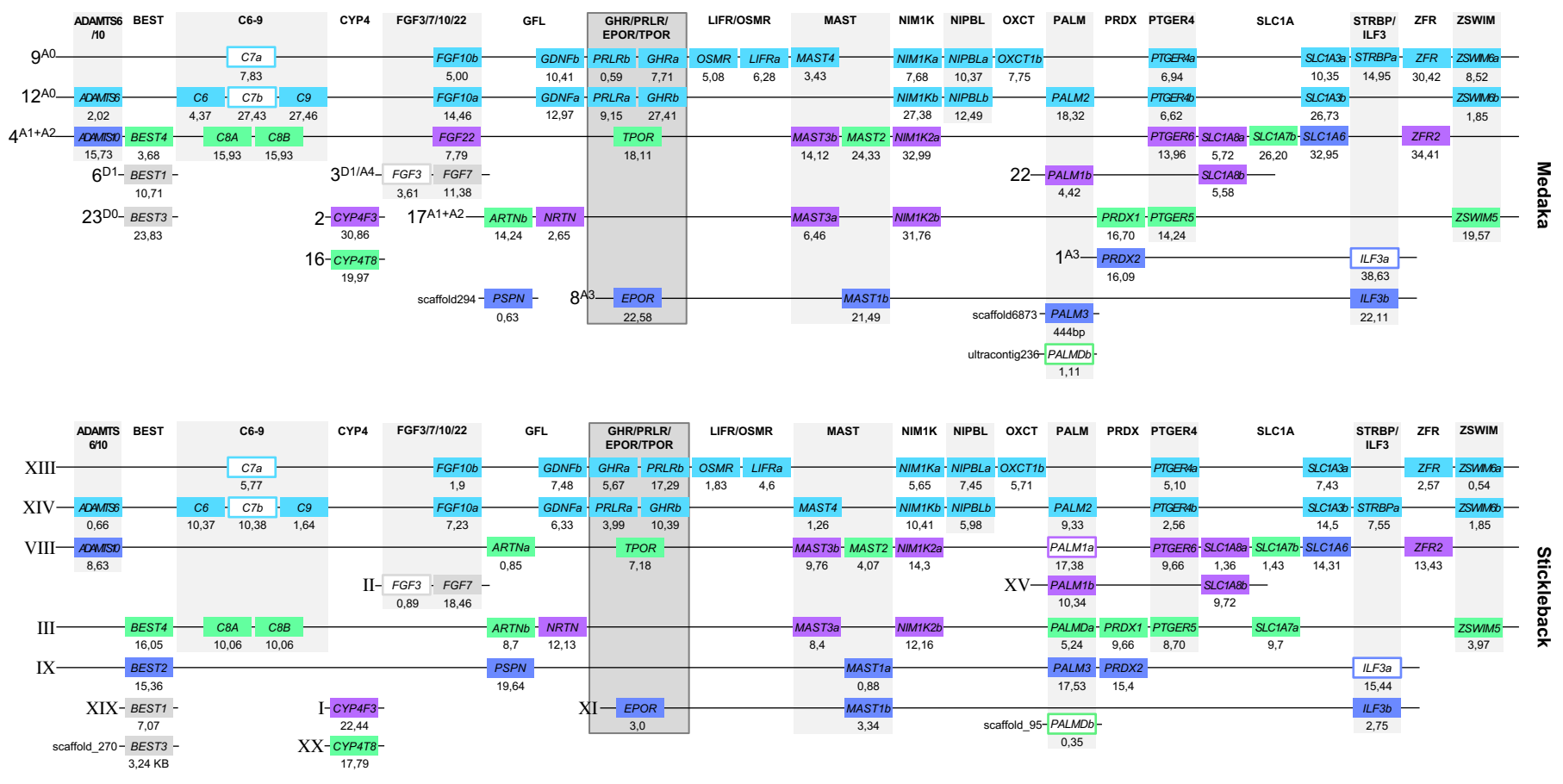
Medaka
ADAMTS6/10
BEST
C6-9
CYP4
FGF3/7/10/22
GFL
GHR/PRLR/EPOR/TPOR
LIFR/OSMR
MAST
NIM1K
NIPBL
OXCT
PALM
PRDX
PTGER4
SLC1A
STRBP/ILF3
ZFR
ZSWIM
9A0
C7a 7,83
FGF10b 5,00
GDNFb 10,41
PRLRb 0,59
GHRa 7,71
OSMR 5,08
LIFRa 6,28
MAST4 3,43
NIM1Ka 7,68
NIPBLa 10,37
OXCT1b 7,75
PTGER4a 6,94
SLC1A3a 10,35
STRBPa 14,95
ZFR 30,42
ZSWIM6a 8,52
12A0
ADAMTS6 2,02
C6 4,37
C7b 27,43
C9 27,46
FGF10a 14,46
GDNFa 12,97
PRLRa 9,15
GHRb 27,41
NIM1Kb 27,38
NIPBLb 12,49
PALM2 18,32
PTGER4b 6,62
SLC1A3b 26,73
ZSWIM6b 1,85
4A1+A2
ADAMTS10 15,73
BEST4 3,68
C8A 15,93
C8B 15,93
FGF22 7,79
TPOR 18,11
MAST3b 14,12
MAST2 24,33
NIM1K2a 32,99
PTGER6 13,96
SLC1A8a 5,72
SLC1A7b 26,20
SLC1A6 32,95
ZFR2 34,41
6D1
BEST1 10,71
3D1/A4
FGF3 3,61
FGF7 11,38
22
PALM1b 4,42
SLC1A8b 5,58
23D0
BEST3 23,83
2
CYP4F3 30,86
17A1+A2
ARTNb 14,24
NRTN 2,65
MAST3a 6,46
NIM1K2b 31,76
PRDX1 16,70
PTGER5 14,24
ZSWIM5 19,57
16
CYP4T8 19,97
1A3
PRDX2 16,09
ILF3a 38,63
scaffold294
PSPN 0,63
8A3
EPOR 22,58
MAST1b 21,49
scaffold6873
PALM3 444bp
ILF3b 22,11
ultracontig236
PALMDb 1,11
Stickleback
ADAMTS 6/10
BEST
C6-9
CYP4
FGF3/7/10/22
GFL
GHR/PRLR/EPOR/TPOR
LIFR/OSMR
MAST
NIM1K
NIPBL
OXCT
PALM
PRDX
PTGER4
SLC1A
STRBP/ILF3
ZFR
ZSWIM
XIII
C7a 5,77
FGF10b 1,9
GDNFb 7,48
GHRa 5,67
PRLRb 17,29
OSMR 1,83
LIFRa 4,6
NIM1Ka 5,65
NIPBLa 7,45
OXCT1b 5,71
PTGER4a 5,10
SLC1A3a 7,43
ZFR 2,57
ZSWIM6a 0,54
XIV
ADAMTS6 0,66
C6 10,37
C7b 10,38
C9 1,64
FGF10a 7,23
GDNFa 6,33
PRLRa 3,99
GHRb 10,39
MAST4 1,26
NIM1Kb 10,41
NIPBLb 5,98
PALM2 9,33
PTGER4b 2,56
SLC1A3b 14,5
STRBPa 7,55
ZSWIM6b 1,85
VIII
ADAMTS10 8,63
ARTNa 0,85
TPOR 7,18
MAST3b 9,76
MAST2 4,07
NIM1K2a 14,3
PALM1a 17,38
PTGER6 9,66
SLC1A8a 1,36
SLC1A7b 1,43
SLC1A6 14,31
ZFR2 13,43
II
FGF3 0,89
FGF7 18,46
XV
PALM1b 10,34
SLC1A8b 9,72
III
BEST4 16,05
C8A 10,06
C8B 10,06
ARTNb 8,7
NRTN 12,13
MAST3a 8,4
NIM1K2b 12,16
PALMDa 5,24
PRDX1 9,66
PTGER5 8,70
SLC1A7a 9,7
ZSWIM5 3,97
IX
BEST2 15,36
PSPN 19,64
MAST1a 0,88
PALM3 17,53
PRDX2 15,4
ILF3a 15,44
XIX
BEST1 7,07
I
CYP4F3 22,44
XI
EPOR 3,0
MAST1b 3,34
ILF3b 2,75
scaffold_270
BEST3 3,24 KB
XX
CYP4T8 17,79
scaffold_95
PALMDb 0,35