

Homo sapiens					
Chr 13	27018052	27092548	-	Lnx2	
Chr 13	27094003	27139547	+	PolR1d	
Chr 13	27264780	27266089	+	Gsh-1	
Chr 13	27392157	27397394	+	Ipf-1(Pdx)	
Chr 13	27434281	27441317	-	Cdx-2	
Chr 13	27475753	27572703	-	Flt3	ENSG00000183463
Chr 13			+	Pan3	
Chr 13	27775307	27966980	-	Flt1 (Vegfr1)	
Chr 13	28131292	28150335	+		AF275807
Chr 13	28986497	29025998	-	SLC7A1	
Chr 13	29933497	29935817	-	Hmg1	
chr X	71332388	71575674	-	Hdac8	
chr X	71583502	71716903	-	Kpb1	
Chr X		72278376	+		ENSG00000185854
Chr X	72450111	72457442	+	Cdx-4	
Chx X	72566065	73128879	+	Chic1	
Chr 4	54121708	54166541	+	Fip1L1	
Chr 4	54166396	54265295	-	Lnx1	
Chr 4	54771716		-	Cchic2	
Chr 4	54807126	54809050	+	Gsh-2	
Chr 4	54936130		+	PdgfrA	
Chr 4	55039987	55063197	+	skoygly	
chr 4	55365089	55447783		Kit	
chr 4	55787039	55832388	-	Kdr (Vegfr2)	
chr 5	149413052	149473128	-	K-Fms (Csf1r)	
Chr 5	149515614		-	PdgfrB	
Chr 5	149526552	149544256	+	Cdx-1	
Chr 5	149549713		+	SLC6A7	
Chr 5	149579248	149649511	-	Camk2A	
Chr 5	179968573	180009151	-	Flt4 (Vegfr3)	
Rattus norvegicus					
Chr 12	7888255	8065404	+	Flt1	
Chr 12	8239722	8296319	+	Flt3	
Chr 12	8326786	8332395	+	Cdx?	ENSRNOESTG00000008932
				<i>rather obvious assembly problems</i>	
Chr 12	8389470	8414416	-	Lnx2	
Chr 12	8452288	8452659	+	Rpa9	
Chr 12	8581078	8582373	+	Gsh-1	
Chr X	90431712	90617611	-	Hdac8	
Chx X	91359230	91367068	+	Cdx-1	ENSRNOG00000002974
Chr X	91382409		+	S26E	
Chr X	91394864	91432607	+	Chic1	
	?	?		Kdr (Vegfr2)	
Chr 14	34906884	34984775	-	Kit	
Chr 14	35372540	35407913	-	PdgfrA	
Chr 14	35409169		-	Pgds	
Chr 14	35489441	35491034	-	Gsh-2	
Chr 14	35522808	35556776	+	Chic2	
Chr 14	35905536	35992947	+	Lnx1	
Chr 14	35994219	36024504	-	Fip1L1	
Chr 18	56951520	57020491	+	Camk2a	
Chr 18	57055516	57076563	-	Cdx-1	
Chr 18	57103107	57124100	+	PdgfrB	
Chr 18	57115842	57177481	+	Fms (Csf1r)	
Chr 10	35087059	35109326	+	Flt4 (Vegfr3)	

Mus musculus					
Chr 5	144342135	144401560	-	Lnx2	
Chr 5		144404111	+	Rpo1-3	
Chr 5	144513345	144514829	+	Gsh-1	
Chr 5	144594959	144600680	+	Ipf-1 (Pdx)	
Chr 5	144625733	144632082	-	Cdx-2	
Chr 5		144656025	-	Flt3	
Chr 5	144887029	145050821	-	Flt1 (Vegfr1)	
Chr 5	145185470	145200613	+	BC027016	
Chr 5	146381837	146383720	-	Hmg1	
Chr X	93711216	93908643	-	Hdac8	
Chr X	94660193		-	Q8K3S0	
Chr X	94724977	94733803	+	Cdx-4	
Chr X	94760082		+	Chic1	
Chr 5	73260794	73292327	+	Fip1L1	
Chr 5	73293448	73399533	-	Lnx1	
Chr 5	73740090		-	Chic2	
Chr 5	73771071	73773363	+	Gsh-2	
Chr 5	73847970		+	PdgfrA	
Chr 5	74270600	74350808	+	Kit	
Chr 5	74630299	74673626	-	Kdr (Vegfr2)	
Chr 18	61439334	61501696	+	Camk2a	
Chr 18	61527897		-	Slc6a7	
Chr 18	61532563	61549897	-	Cdx-1	
Chr 18	61559139	61598757	+	PdgfrB	
Chr 18	61619575	61644835	+	K-Fms (Csf1r)	
Chr 11	49264383	49290819	+	Flt4 (Vegfr3)	
Canis familiaris					
Chr 25	12538140	12540592	+	Hmg1	
Chr 25	14217669	14392845	+	Flt1	
Chr 25	14602884	14648310	+	Flt3	
Chr 25	14687192	14693175	+	Cdx-2	ENSCAFG00000006742
Chr 25	14728543	14732931	-	Ipf-1	ENSCAFG00000006746
Chr 25	14840995	14842258	-	Gsh-1	ENSCAFG00000006749
Chr 25	14984227	14984601	-	PolR1d	ENSCAFG00000006755
Chr 25	15026492	15061458	+	Lnx2	
Chr X	59247952	59434250	-	Hdac8	
Chr X	60071799		-	Nap1l2	
Chr X	60107366	60118535	+	Cdx-4	
Chr X	60121699			UNK	GENSCAN00000017930
Chr X	60182347	60237535	+	Chic1	
Chr 13	49164568	49204822	+	Fip1L1	
Chr 13	49206245	49308808	-	Lnx1	
Chr 13	49682051	49728780	-	Chic2	
Chr 13	49763459	49763632	+	Gsh-2	ENSCAFG00000002052
Chr 13	49861528	49894904	+	PdgfrA	
Chr 13	49879380	50308953	+	Kit/Flt3/?	
Chr 13	50564101	50601558	-	Kdr (Vegfr2)	
Chr 4	61969395	62030707	+	Camk2a	
Chr 4	62055002		-	SLC6A7	ENSCAFG00000018203
Chr 4	62067272	62084179	-	Cdx-1	ENSCAFG00000018210
Chr 4	62094911	62176613	+	PdgfrB	NP_001003382
Chr 4	62122146	62176616	+	Fms (Csf1r)	
Chr 11	10793511	10826868	-	Flt4 (Vegfr3)	

Monodelphis domestica					
scaf_120	445937	627670	+	Vegfr1/?	
scaf_13183	454775	456700	-	Flt3	(824,79%)
scaf_13292	79577	119834	+	Flt3	(1103,88%)
scaf_13292	156740	162492	+	Cdx-2	
scaf_13292	201674	201841	-	Cdx-1/?	
scaf_13292	352793	353921	-	Gsh-1	
scaf_13292	608927	643513	+	Lnx2	
scaf_1784	22473	25447	-	Cdx-4/?	
scaf_1784	54086	59605	-	Cdx-2/?	
scaf_1046	84392	115935	+	Kdr/?	
scaf_18580	4804260	4846049	+	Kdr	(2558,86%)
scaf_18580	5155356	5201980	-	Kit	
scaf_18580	5734698	5765846	-	PdgfrA	
scaf_18580	5885662	5885847	-	Gsh-2/?	
scaf_18580	5935314	5984651	+	Chic2	
scaf_18580	6525634	6660958	+	Lnx1	
scaf_18580	6662488	6724135	-	Fip1L1	
scaf_15003	2661935	2754444	-	Fms (Csf1r)	
scaf_15003	2742098	2782487	-	PdgfrB	
scaf_15003	2871172	2872876	+	Cdx-1/?	
scaf_15003	2937355	3017816	-	Camk2a	
scaf_18574	3099988	3155134	+	Vegfr3	(2109,86%)
Gallus gallus					
Chr 1	167864641	167941388	+	Flt1	
Chr 1	168067062	168070960	+	Flt3	
Chr 1	168079835	168082614	+	Cdx-2/?	
Chr 1	168096505	168096846	-	Ipf-1 (Pdx)	
Chr 1	168125596	168125796	-	Gsh-1	
Chr 1		168161724	-	PolR1d	ENSGALG00000017095
Chr 1	168212578	168237000	+	Lnx2	
Chr 4	11764577	11781479	+	Flt1/?	
Chr 4	11838514	11850537	+	Cad1 (Cdx-4)	ENSGALG00000007656
Chr 4	11859301	11867657	+	Cad1 (Cdx-4)	
Chr 4	11880390	11890176	+	Chic1	
Chr 4	11923410	11939297	-	Lnx2	
Chr 4	65667038	65691708	+	Kdr (Vegfr2)	
Chr 4	65752516	65802420	-	Kit	
Chr 4	65879695	65902450	-	PdgfrA	
Chr 4	65941675	65941899	-	Gsh-2 (part)	
Chr 4	65954455	65975378	+	Chic2	
Chr 4	66137839	66198904	+	Lnx1	
Chr 4	66200458	66217038	-	Fip1L1	
Chr Un	97586045	97586380	-	Gsh-2/?	XM_427550
Chr 13	12133573	12151964	+	Camk2a	
Chr 13	12173782	12174131	-	Cdx-2/?	
Chr 13	12165366		-	SLC6A7	
Chr 13	12172161	12182384	-	Cdx-1/?	
Chr 13	12201604	12208349	+	PdgfrB	ENSGALT00000009189
Chr 13	12227385	12230534	+	Fms (Csf1r)	ENSGALT00000009190
Chr 13	12321110	12325701	+	Flt4(Vegfr3)	ENSGALT00000009188

Xenopus tropicalis				
scaf_449	392664	399738	-	Lnx2
scaf_449	558244	560262	+	Gsh-2/?
scaf_449	611644	634265	+	Pdx
scaf_449	654925	656036	-	Cdx-2/?
scaf_449	718979	728200	-	Flt3
scaf_449		875241	-	Flt1
scaf_69	2275525	2288553	+	Lnx2'
scaf_69	2314608	2321759	-	Chic1
scaf_69	2326751	2327241	-	Cdx-4
scaf_69	2387671	2392835	-	Kdr (Vegfr2)
scaf_69	2390912	2404587	-	Flt1/?
scaf_248	519754	546830	+	Fip1L1
scaf_248	551579	642917	-	Lnx1
scaf_248	774542	792448	-	Chic2
scaf_248	808887	810158	+	Gsh-2
scaf_248	855484	868411	+	PdgfrA
scaf_248	1004798	1011924	+	Kit
scaf_248	1103987	1110991	-	Kdr (Vegfr2)
scaf_635	341854	443555	+	Camk2a
scaf_635	514203		-	SLC6A2
scaf_635	528842	544328	-	Cdx-1
scaf_635	550072			GENSCAN00000103921
scaf_635	587808			GENSCAN00000103928
scaf_635	619751	648818	+	PdgfrB
scaf_635	692871	698817	+	Fms (Csf1r)
scaf_10	259778	294590	-	Flt4 (Vegfr3)
Takifugu rubripes				
Scaff_146	51084	68453	+	Flt1
Scaff_146	82687	84594	+	Flt3
Scaff_146	86907	88902	-	Ipf-1 (Pdx)
Scaff_146	95479	101000	+	Lnx2
Scaff_6		394619	+	SINFRUG00000149943
Scaff_6	404571	405414	-	Gsh-1
Scaff_6	427424		+	SINFRUG00000149948
Scaff_580		- 47179	+	arrestin-b2
Scaff_580	51644	52142	+	Cdx/?
Scaff_580	65692		+	Ins
Scaff_598	57721	- 80291	+	Kdr
Scaff_598	87649	90172	+	Cad1 (Cdx)
Scaff_598	91236	94584	+	Chic1
Scaff_598	97149	102841	-	Lnx/?
Scaff_973	67205	67558	-	Lnx1
Scaff_827	end			
Scaff_827	6516	7402	+	Gsh-2
Scaff_827	8984	19284	+	PdgfrA
Scaff_827	32491	48672	+	Kit
Scaff_827	67844	77106	-	Kdr
Scaff_4094	2762	4396		Chic2
Scaff_1324	23298	30370	-	Csf1r
Scaff_1324	32730	43177	-	PdgfrB
Scaff_1324	55201	57951	+	Cdx-1
Scaff_1324	end			
Scaff_663	118842	129794	-	Flt4
Scaff_525	52541	58079	+	PdgfrB2
Scaff_525	61723	67394	+	Csf1r2

Oryzias latipes				
scaff_3770	37044	37409	-	Lnx/?
scaff_3770	44702	46914	+	Pdx
scaff_602	82438	83272	+	Gsh-1
scaff_18138	3016	3474	+	Cdx2/1/?
scaff_1170	2903	3343	+	Lnx2
scaff_1170	13815	13657	-	chic/?
scaff_1170	18154	20471	-	Cdx-4
scaff_1170	31297	69603	-	Flk1/Vegfr2/?
scaff_7663	8450	9330	+	Gsh-2
scaff_7663	19386	29149	+	PdgfrA
scaff_1722	43048	60300	-	Kdr
scaff_692	34006	38222	-	Fms (Csf1r)
scaff_692	42636	54509	-	PdgfrB
scaff_692	74427	79787	+	Cdx-1
Danio rerio				
Chr 24	16459561	16465929	-	Flt3/?
Chr 24	16521480	16528326	-	Flt1
Chr 24	16815491	16840431	-	Lnx2
Chr 24	16852534	16854674	+	Ipf-1 (Pdx)
Chr 5	957639	957806	-	Gsh/?
Chr 5	67475253		-	ENSDARG00000026730
Chr 5	67535641	67536461	+	Gsh-1
Chr 5	67553857		+	ENSDARG00000026739
Chr 14	47608491	47669829	+	Kdr (Vegfr2)
Chr 14	47641981	47648504	+	Vegfr/?
Chr 14	47690116	47692885	+	Cdx-4
Chr 14	47697317	47702099	+	Chic1
Chr 14	47709946	47809210	-	Lnx2/?
				<i>rather obvious assembly problems</i>
Chr 14	47737324	47744215	+	Lnx2/?
Chr 14	47752083	47756881	-	Chic1
Chr 14	47761298	47764013	-	Cdx-4
Chr 14	47804211	47804860	-	Kdr (Vegfr2)
Chr 20	26351293	26355554	-	Vegfr1/Kdr/?
Chr 20	26420356	26462629	-	Kit
Chr 20	26521191	26548346	-	PdgfrA
Chr 20	26567796	26568825	-	Gsh-2
Chr 20	26765367	27200068	+	Lnx1
				<i>rather obvious assembly problems</i>
Chr 20	27146596		-	PdgfrA/?
Chr 20	27158291	27159320	-	Gsh-2
Chr 20	27179116		+	ENSDARG00000031803
Chr 20	27210083	27210283	+	Lnx1
Chr 20	27222619	27269655	-	Fip1L1
Chr 2	20446456	20496442	-	Csfr1
Chr 2	20455858	20456441	-	pdgfrB
Chr 23	41466007	41480005	-	Flt4/? (Vegfr3)
Chr NA	2668531	2668899	-	Cdx-1a

Tetraodon nigroviridis					
Chr 6	2683552	2686275	+	Flt1	
Chr 6	2706985	2708313	+	Flt3	
Chr 6	2710900	2712765	-	Ipf-1 (Pdx)	
Chr 6		2714223	-	RNApol or	
Chr 6	2716250	2723445	+	Ln timer	
Chr 7		4243646	+		GSTENG00019039001
Chr 7	4250981	4251857	+	Gsh-1	
Chr 7		4258099	-		GSTENG00019043001
Un_random	162747406	162747531	-	Cdx4/?	
Un_random	68186467	68186859	-	Cdx4/?	
Chr 1	8363701	8370648	-	Kdr/?	
l_random	1979905	1999041	-	Ln timer	
l_random	2039772		-		
l_random	2049256	2049874	+	Gsh-2	
l_random	2057599	2067439	+	PdgfrA	
l_random	2088534	2092113	+	Kit	
l_random	2113139	2114247	-	Kdr (Vegfr2)	
Chr 18	3400312	3407996	-	Kit	
Chr 18	3418931	3425408	+	Fip1L1	
Chr 18	3417815	3417934	+		
Chr 18	3439152	3450889	+	Chic2	
Chr 1		413879	+	Arylsulfatase	GSTENG00026574001
Chr 1	424038	424187	-	Cdx-1/?	GIDT00020444001
Chr 1	444812	446524	-	Cdx-1	GSTENG00026576001
Chr 1	458257	466985	+	PdgfrB	
Chr 1	471619	475803	+	Fms (Csf1r)	
Chr 1	2079073	2081633	-	Flt4/? (vegfr3)	
Chr 7	9994999	9999310	+	Cdx-1/?	
Chr 7	10009603	10010614	+	Ins	
Chr 7	10883736	10885040	-	Fms (Csf1r)	
Chr 7	10970776	10977555	-	Camk2a	