
Supplementary information

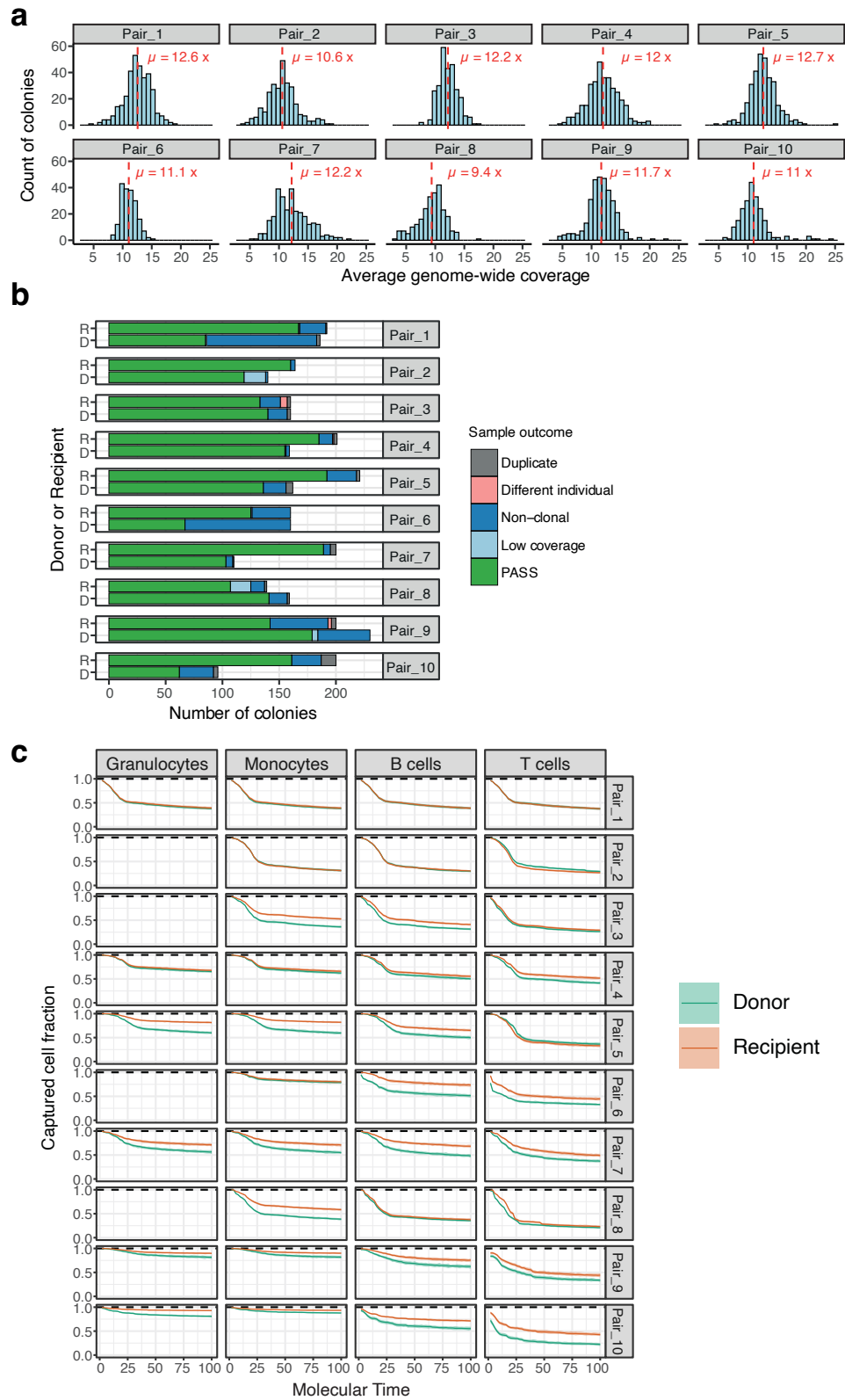
**Clonal dynamics after allogeneic
haematopoietic cell transplantation**

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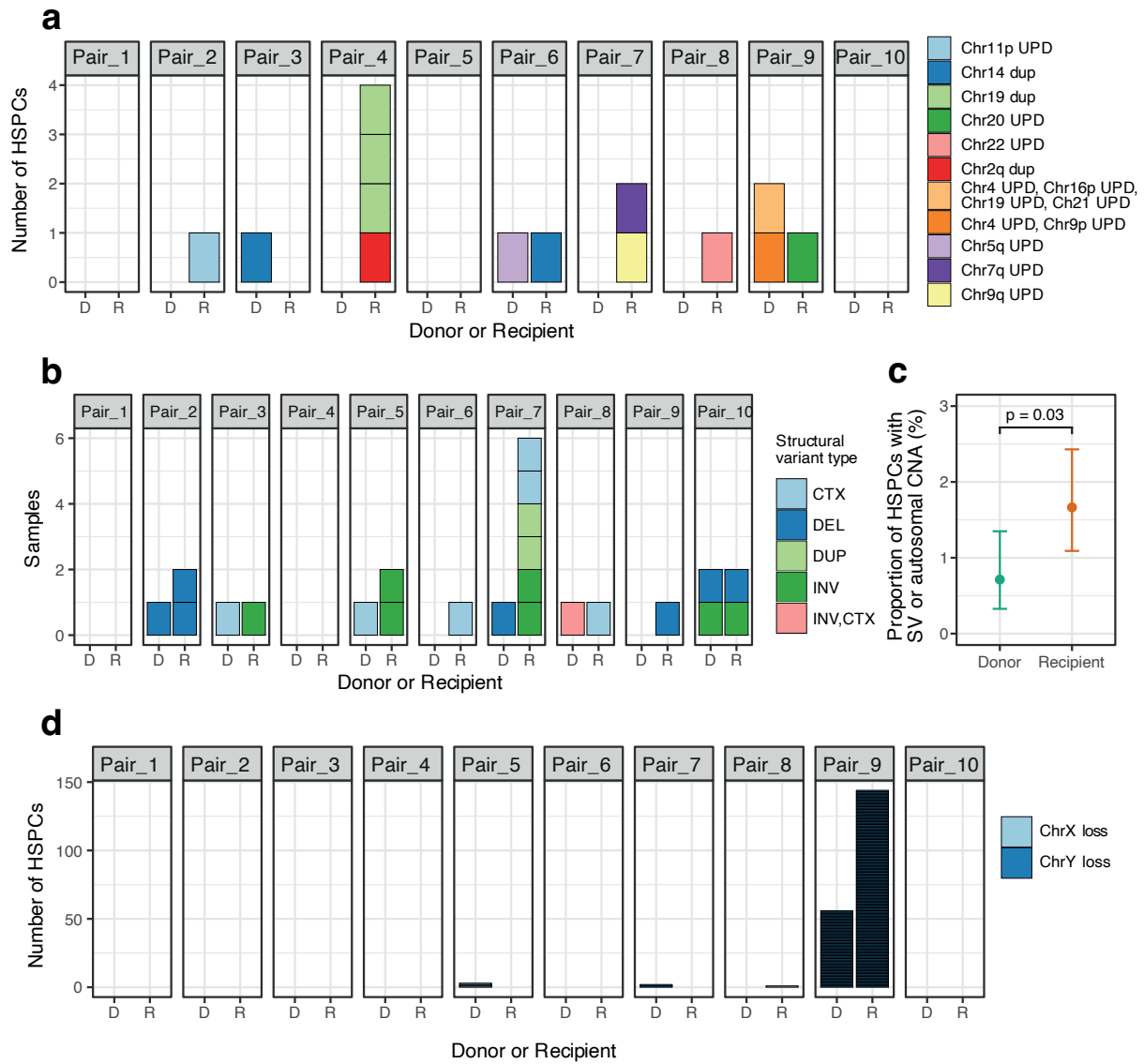
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Clonal dynamics after allogeneic haematopoietic cell transplantation

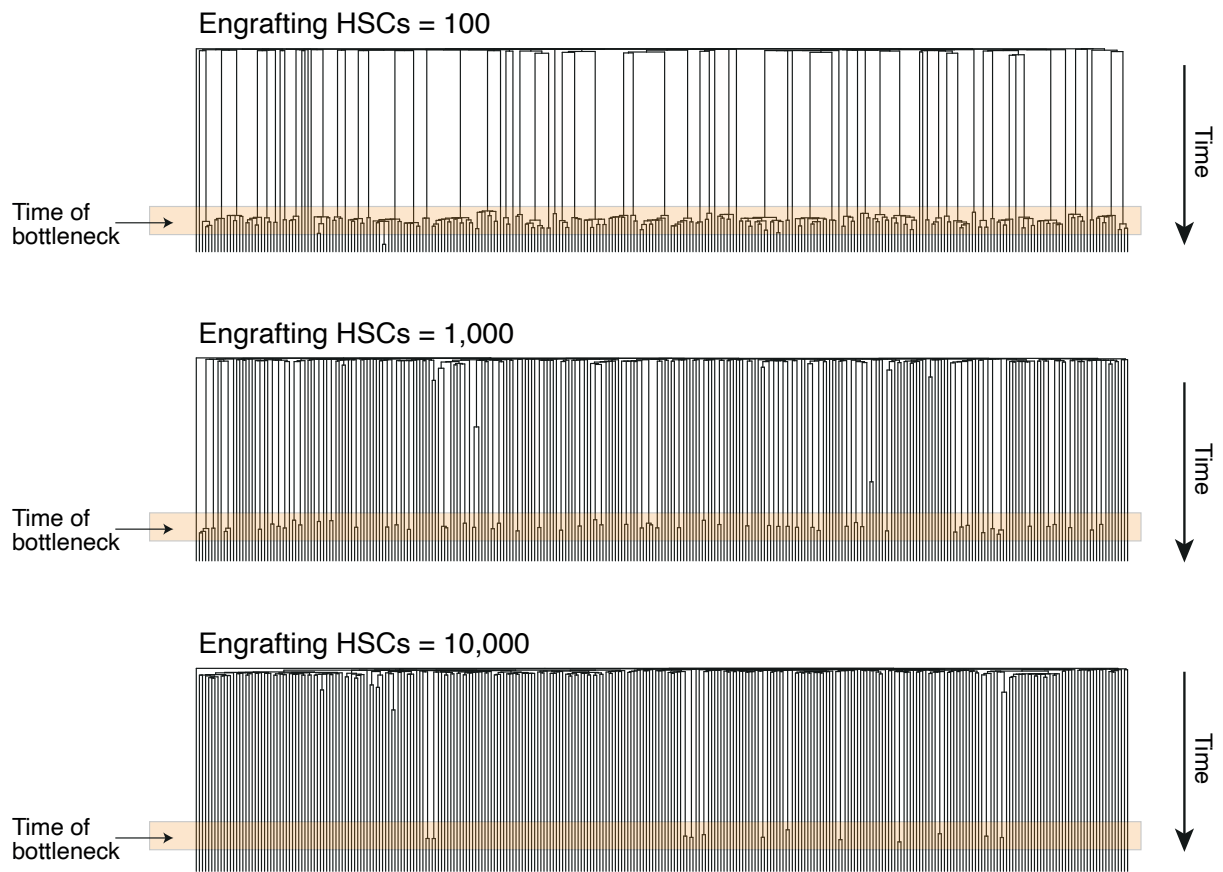
Authors: Michael Spencer Chapman^{1,2,3}, C. Matthias Wilk⁴, Steffen Boettcher⁴, Emily Mitchell^{1,2,3}, Kevin Dawson¹, Nicholas Williams¹, Jan Müller⁴, Larisa Kovtonyuk⁴, Hyunchul Jung¹, Francisco Caiado⁴, Kirsty Roberts¹, Laura O'Neill¹, David G Kent^{2,3,5}, Anthony R. Green^{2,3}, Jyoti Nangalia^{1,2,3}, Markus G. Manz⁴ *, Peter J. Campbell^{1,2,3} *



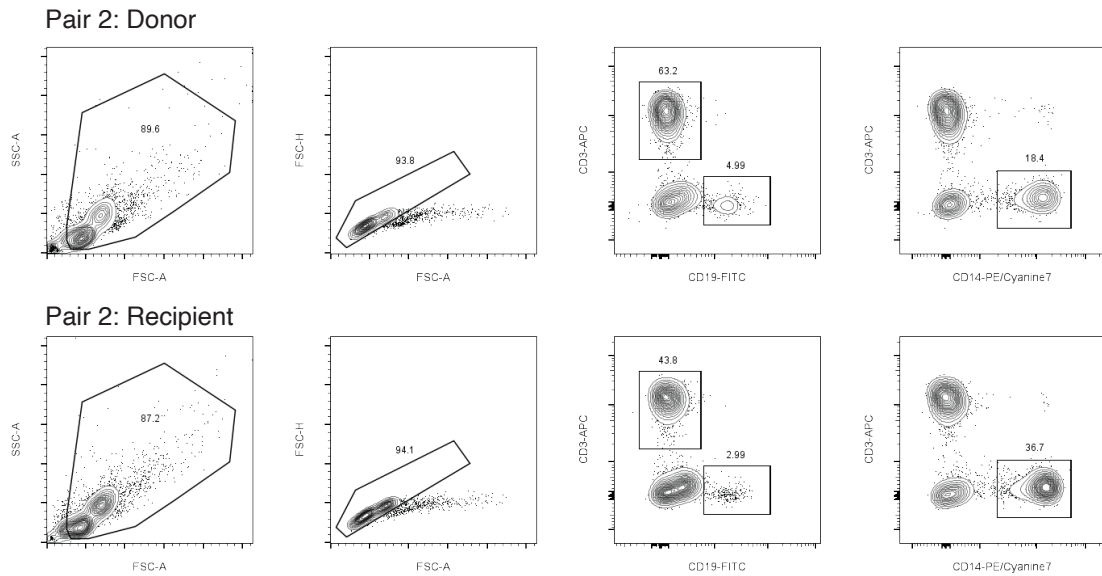
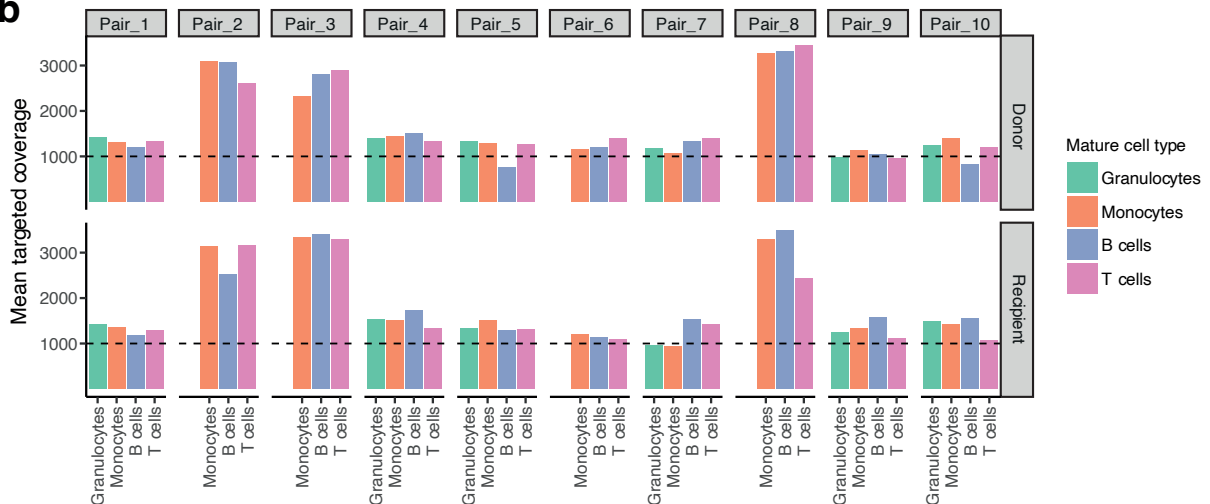
Supplementary Fig. 1 | Sequencing metrics and quality control. **a**, Histograms of the mean whole genome sequencing coverage of colonies used in the final data set. Red dashed lines indicate the mean value across an individual, which is also printed on each panel. **b**, Bar plot showing the outcome of all colonies undergoing WGS, divided by sibling transplant pair and donor or recipient status. **c**, Line plot showing the sum of clonal fractions across the branches of the phylogenetic tree at different points in molecular time. This is divided by pair and by cell type. The earliest time point shows the sum of clonal contributions of the first two blastomeres of the embryo. Solid line shows the median posterior values, shaded areas show the 95% posterior intervals.



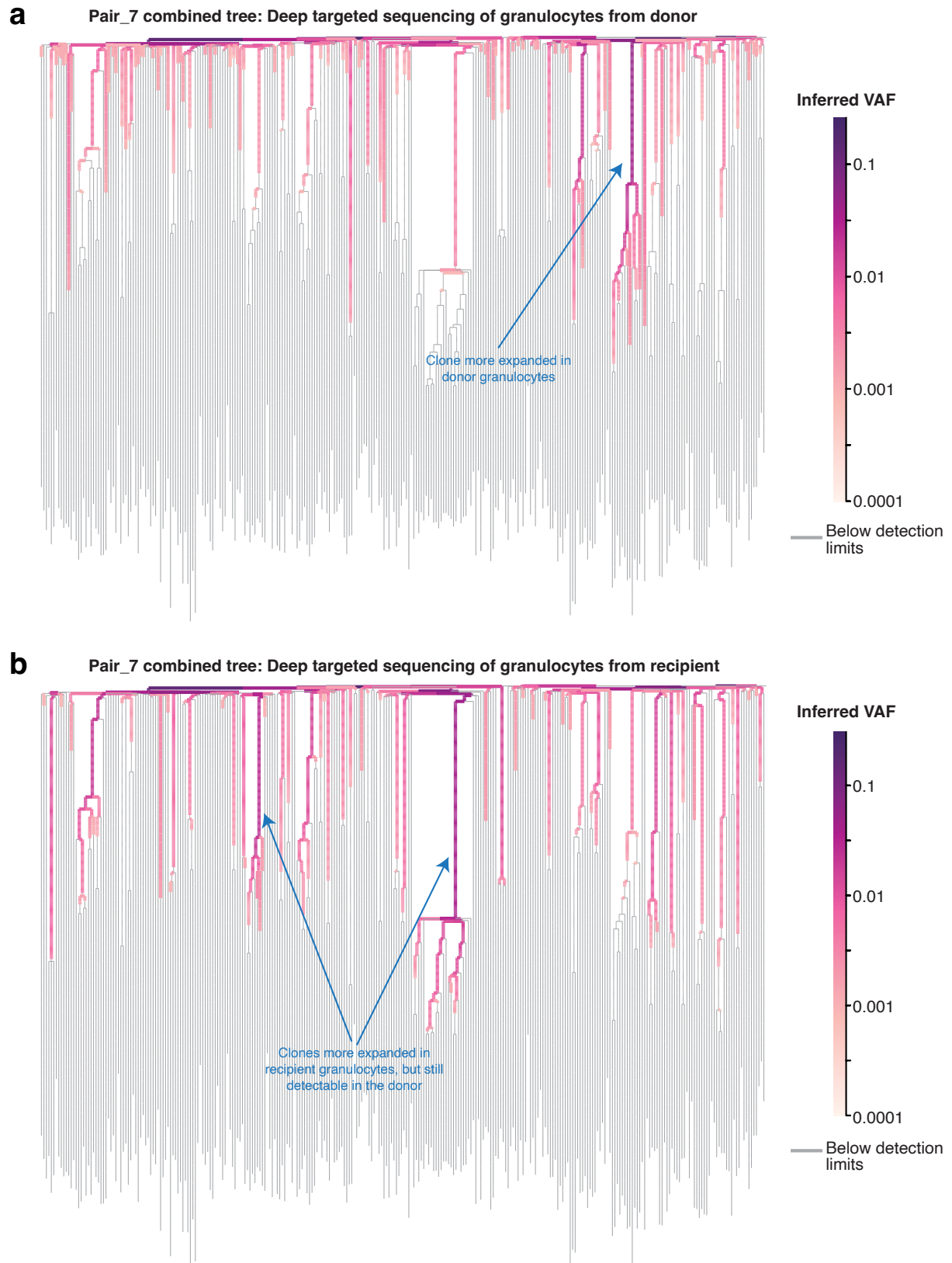
Supplementary Fig. 2 | Copy number alterations and structural variants. **a**, Stacked bar plot showing the number of HSPCs with autosomal CNAs in each individual, coloured by CNA type. **b**, Stacked bar plot showing the number of HSPCs with structural variants (SVs) in each individual, coloured by the class of SV. **c**, Dot plot showing the proportion of HSPCs harbouring either a CNA or SV combined across all HCT donors (green) and recipients (orange). Error bars show the 95% confidence interval (binomial test) with the p value calculated using Pearson's chi-square test. **d**, As in **a**, but for CNAs in the X and Y chromosomes. UPD, Uniparental disomy; dup, duplication; del, deletion; CTX, reciprocal translocation; INV, inversion; INV/ CTX, complex inversion and translocation event.



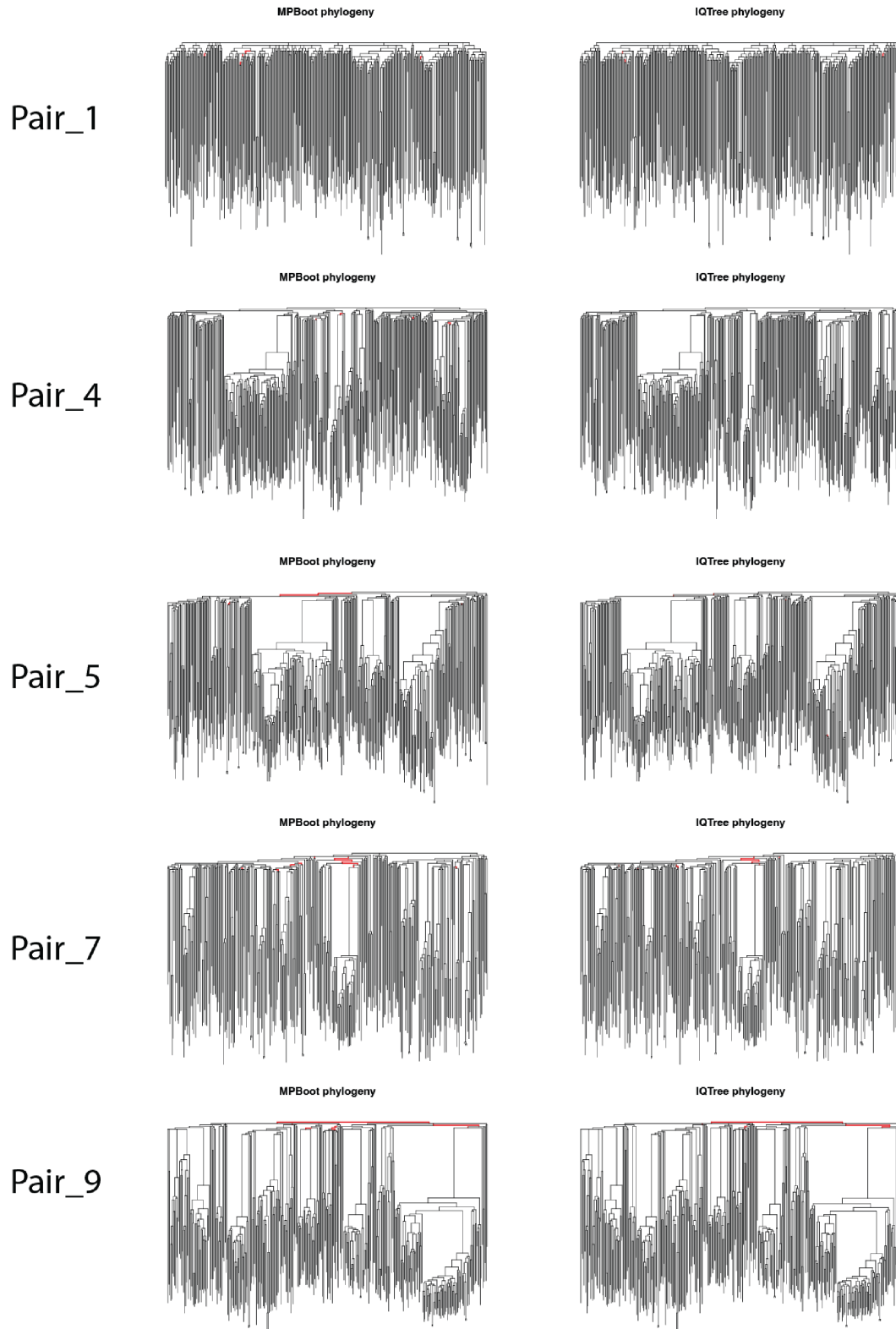
Supplementary Fig. 3 | Bottleneck simulations. Simulations of a transplant bottleneck of varying degrees. Left - bottleneck of 100 HSCs, middle - bottleneck of 1000 HSCs, right - bottleneck of 10,000 HSCs. The orange box highlights the time of the bottleneck. See methods for details of simulation framework.

a**b**

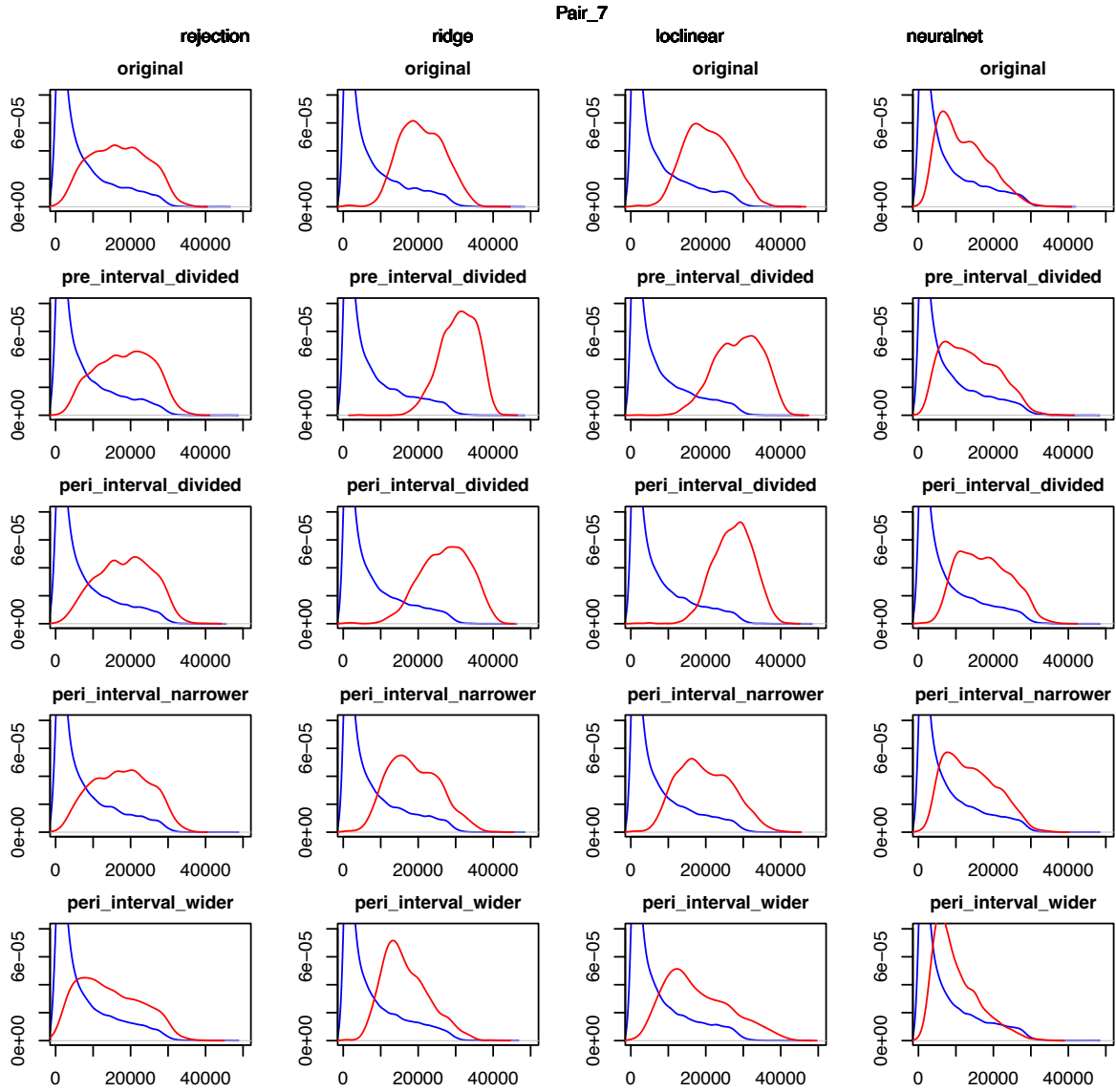
Supplementary Fig. 4 | Deep targeted sequencing of flow-sorted mature cell populations. a, Gating strategy for flow cytometric cell sorting of B-cells, T-cells and monocytes from peripheral blood cells, as illustrated by the gating used for Donor and Recipient Pair 2. **b**, Mean target sequencing coverage of bulk samples undergoing targeted sequencing, using a bait-capture approach. Results are divided by individual and cell type. Eight of 20 individuals did not have granulocytes available for sequencing. The horizontal dashed line shows the 1000x coverage value - the minimum targeted in these experiments.



Supplementary Fig. 5 | Bayesian inference of variant allele fractions from deep targeted sequencing. a, Combined phylogeny for Pair_7, with mutations coloured by the estimated posterior median VAF from deep targeted sequencing data on a sample of granulocytes from the donor. **b,** as in **a**, but for a sample of granulocytes from the recipient.



Supplementary Fig. 6 | Comparison of five phylogenies generated by MPBoot with those generated by IQtree. These are the raw phylogenies without branch length correction or removal of duplicates. Short branches with <10 mutations assigned have been extended to a length of 10 to facilitate visualisation of differences. Branches present in one tree but absent in the other are highlighted in red.



Supplementary Fig. 8 | Robustness of the posterior distributions to different definitions of epochs used in the summary statistics. Each facet plot shows the prior (blue) and estimated posterior (red) distributions for size of the HSCT bottleneck for one of the donor-recipient pairs (Pair 7). From left to right, the facets are organised by different regression methods for estimating the posterior parameter values within the ABC. From top to bottom, the facets are organised by the different definitions of epochs (original method on top followed by the four different definitions described above).

Supplementary Table 1 | List of genes assessed for possible driver mutations in haematopoietic stem cell transplant donor-recipient pairs.

<i>ABL1</i>	<i>ALK</i>	<i>ARID1A</i>	<i>ARID2</i>	<i>ASXL1</i>	<i>ASXL2</i>
<i>ATR</i>	<i>BAP1</i>	<i>BAX</i>	<i>BCL10</i>	<i>BCL2</i>	<i>BCOR</i>
<i>BCORL1</i>	<i>BRAF</i>	<i>BRCC3</i>	<i>CALR</i>	<i>CBL</i>	<i>CBLB</i>
<i>CBLC</i>	<i>CCDC115</i>	<i>CCL22</i>	<i>CDK4</i>	<i>CDKN1B</i>	<i>CDKN2A</i>
<i>CDKN2B</i>	<i>CDKN2C</i>	<i>CEBPA</i>	<i>CHEK2</i>	<i>CREBBP</i>	<i>CRLF2</i>
<i>CSF1R</i>	<i>CSF3R</i>	<i>CTCF</i>	<i>CTNNB1</i>	<i>CUX1</i>	<i>DICER1</i>
<i>DNMT3A</i>	<i>DNMT3B</i>	<i>DUSP22</i>	<i>EED</i>	<i>EGFR</i>	<i>EP300</i>
<i>ETV6</i>	<i>EZH2</i>	<i>FAM175A</i>	<i>FBXW7</i>	<i>FGFR2</i>	<i>FLT3</i>
<i>GATA1</i>	<i>GATA2</i>	<i>GNAS</i>	<i>GNB1</i>	<i>H3F3A</i>	<i>H3F3B</i>
<i>HRAS</i>	<i>IDH1</i>	<i>IDH2</i>	<i>IGLL5</i>	<i>IKZF1</i>	<i>IRF4</i>
<i>JAK2</i>	<i>JAK3</i>	<i>KDM5C</i>	<i>KDM6A</i>	<i>KIT</i>	<i>KMT2A</i>
<i>KRAS</i>	<i>MAGEC3</i>	<i>MGA</i>	<i>MPL</i>	<i>MTA2</i>	<i>MYC</i>
<i>MYD88</i>	<i>NF1</i>	<i>NF2</i>	<i>NFE2L2</i>	<i>NOTCH1</i>	<i>NOTCH2</i>
<i>NPM1</i>	<i>NRAS</i>	<i>PAX5</i>	<i>PDGFRA</i>	<i>PHF6</i>	<i>PHIP</i>
<i>PIK3CA</i>	<i>PPM1D</i>	<i>PTEN</i>	<i>PTPN11</i>	<i>RAC1</i>	<i>RAD21</i>
<i>RAD50</i>	<i>RAD51</i>	<i>RB1</i>	<i>RHOA</i>	<i>RRAS</i>	<i>RUNX1</i>
<i>SETBP1</i>	<i>SETD2</i>	<i>SF3B1</i>	<i>SH2B3</i>	<i>SIK3</i>	<i>SMC1A</i>
<i>SMC3</i>	<i>SPRED2</i>	<i>SRCAP</i>	<i>SRSF1</i>	<i>SRSF2</i>	<i>STAG2</i>
<i>STAT3</i>	<i>STAT5A</i>	<i>SUZ12</i>	<i>TERT</i>	<i>TET2</i>	<i>TP53</i>
<i>U2AF1</i>	<i>WHSC1</i>	<i>WT1</i>	<i>YLPM1</i>	<i>ZBTB33</i>	<i>ZNF234</i>
<i>ZNF318</i>	<i>ZRSR2</i>				

Supplementary Table 2 | Assessment of possible driver mutations in haematopoietic stem cell transplant donor-recipient pairs. The mutation reference is given as chromosome-position-reference-variant allele. The decision column gives the assessment of whether the variant was considered a driver or variant of unknown significance (VUS).

Mutation reference	Node	Gene	Protein change	Pair	Decision
15-90633818-C-T	2	IDH2	IDH2 p.R89H	Pair11	VUS
16-3819192-C-T	130	CREBBP	CREBBP p.G1015R	Pair11	VUS
17-29557401-G-A	449	NF1	NF1 p.?	Pair11	VUS
17-29559129-T-C	404	NF1	NF1 p.L1079P	Pair11	VUS
17-7577120-C-T	16	TP53	TP53 p.R273H	Pair11	Driver
19-17950455-A-T	147	JAK3	JAK3 p.D424E	Pair11	VUS
22-41573996-A-G	174	EP300	EP300 p.N2094S	Pair11	VUS
2-25466800-G-A	324	DNMT3A	DNMT3A p.R635W	Pair11	Driver
3-105470425-G-A	63	CBLB	CBLB p.H202Y	Pair11	VUS
3-105572309-C-T	51	CBLB	CBLB p.R123K	Pair11	VUS
4-106180795-G-A	577	TET2	TET2 p.G1275R	Pair11	Driver
X-119387292-T-G	131	ZBTB33	ZBTB33 p.S8A	Pair11	VUS
X-140984370-G-A	256	MAGEC3	MAGEC3 p.?	Pair11	VUS
X-39913297-T-C	63	BCOR	BCOR p.?	Pair11	Driver
10-112361839-G-A	214	SMC3	SMC3 p.R1003H	Pair13	VUS
11-118374282-A-G	365	KMT2A	KMT2A p.T2559A	Pair13	VUS
11-32450128-G-C	115	WT1	WT1 p.Y228*	Pair13	Driver
13-28644741-A-G	66	FLT3	FLT3 p.S18P	Pair13	VUS
14-95569687-T-C	132	DICER1	DICER1 p.K1349R	Pair13	VUS
16-30734292-A-G	31	SRCAP	SRCAP p.T1301A	Pair13	VUS
16-67650648-G-A	193	CTCF	CTCF p.G318D	Pair13	VUS
17-58740653-C-T	542	PPM1D	PPM1D p.Q520*	Pair13	Driver
17-7578457-C-T	35	TP53	TP53 p.R158H	Pair13	Driver
2-25462020-C-T	485	DNMT3A	DNMT3A p.G796D	Pair13	Driver
2-25982451-T-C	266	ASXL2	ASXL2 p.N280S	Pair13	VUS
20-31024242-C-T	529	ASXL1	ASXL1 p.Q1243*	Pair13	Driver
22-23237636-G-T	165	IGLL5	IGLL5 p.C136F	Pair13	VUS
22-41545841-G-A	252	EP300	EP300 p.C819Y	Pair13	VUS
4-84406209-G-T	137	FAM175A	FAM175A p.T6K	Pair13	VUS
6-43322772-G-C	187	ZNF318	ZNF318 p.S767C	Pair13	VUS
7-101840562-G-A	76	CUX1	CUX1 p.R635H	Pair13	Driver
7-50468212-T-G	71	IKZF1	IKZF1 p.C441G	Pair13	VUS
9-133760615-C-T	17	ABL1	ABL1 p.P999S	Pair13	VUS
9-139402495-T-C	268	NOTCH1	NOTCH1 p.Y1141C	Pair13	VUS
X-119388946-T-C	234	ZBTB33	ZBTB33 p.F559S	Pair13	VUS
X-53441799-T-C	198	SMC1A	SMC1A p.I107V	Pair13	VUS
6-79655862-CG-C	487	PHIP	PHIP p.N1495fs*19	Pair13	Driver
1-120460347-C-T	302	NOTCH2	NOTCH2 p.V1990M	Pair21	VUS
1-27101603-C-T	5	ARID1A	ARID1A p.P1629S	Pair21	VUS
10-112352918-C-A	357	SMC3	SMC3 p.R634S	Pair21	VUS
11-116730042-G-T	307	SIK3	SIK3 p.L796I	Pair21	VUS
19-13054349-A-G	357	CALR	CALR p.?	Pair21	VUS
2-198273204-A-C	299	SF3B1	SF3B1 p.S336A	Pair21	VUS
2-25457177-G-A	566	DNMT3A	DNMT3A p.P904S	Pair21	Driver

2-25458601-T-A	381	DNMT3A	DNMT3A p.I858F	Pair21	Driver
2-25459851-T-C	196	DNMT3A	DNMT3A p.D811G	Pair21	Driver
2-25463541-G-A	509	DNMT3A	DNMT3A p.S714F	Pair21	Driver
2-25466766-C-T	59	DNMT3A	DNMT3A p.?	Pair21	Driver
2-25469965-G-T	357	DNMT3A	DNMT3A p.Y359*	Pair21	Driver
2-25470497-C-G	506	DNMT3A	DNMT3A p.R326P	Pair21	Driver
22-41533746-G-A	601	EP300	EP300 p.W571*	Pair21	Driver
3-47147587-C-T	158	SETD2	SETD2 p.C1580Y	Pair21	VUS
4-106156005-T-A	252	TET2	TET2 p.D302E	Pair21	Driver
4-106164017-G-A	269	TET2	TET2 p.R1176K	Pair21	Driver
4-55129867-C-T	173	PDGFRA	PDGFRA p.T134M	Pair21	VUS
7-140453140-T-C	65	BRAF	BRAF p.T599A	Pair21	Driver
9-37014993-C-T	243	PAX5	PAX5 p.?	Pair21	VUS
X-133549118-G-A	326	PHF6	PHF6 p.V268I	Pair21	Driver
X-39933871-C-T	528	BCOR	BCOR p.R243H	Pair21	Driver
2-25463235-CAGA-C	77	DNMT3A	DNMT3A p.F752delF	Pair21	Driver
X-15841219-CG-C	34	ZRSR2	ZRSR2 p.G436fs*>53	Pair21	Driver
1-120491683-T-C	82	NOTCH2	NOTCH2 p.K849R	Pair24	VUS
10-123310935-G-A	116	FGFR2	FGFR2 p.R165W	Pair24	VUS
12-111885134-G-A	337	SH2B3	SH2B3 p.G341D	Pair24	VUS
12-12871080-G-T	23	CDKN1B	CDKN1B p.A103S	Pair24	VUS
12-58145290-C-T	288	CDK4	CDK4 p.V71I	Pair24	VUS
17-29667647-G-A	275	NF1	NF1 p.R2349H	Pair24	VUS
17-40457631-C-A	174	STAT5A	STAT5A p.L462M	Pair24	VUS
19-33793139-G-T	6	CEBPA	CEBPA p.S61Y	Pair24	Driver
2-25457242-C-T	103	DNMT3A	DNMT3A p.R882H	Pair24	Driver
2-25463248-G-T	302	DNMT3A	DNMT3A p.R749S	Pair24	Driver
2-25463291-G-C	123	DNMT3A	DNMT3A p.F734L	Pair24	Driver
2-25467478-T-C	236	DNMT3A	DNMT3A p.Y533C	Pair24	Driver
20-31016179-G-A	258	ASXL1	ASXL1 p.S142N	Pair24	VUS
20-57415846-G-A	26	GNAS	GNAS p.A229T	Pair24	VUS
22-41523536-C-T	7	EP300	EP300 p.P318S	Pair24	VUS
5-1264660-C-T	180	TERT	TERT p.R901Q	Pair24	VUS
5-1293700-G-A	40	TERT	TERT p.S434F	Pair24	VUS
6-394906-T-C	49	IRF4	IRF4 p.L101S	Pair24	VUS
X-119389023-C-T	239	ZBTB33	ZBTB33 p.R585C	Pair24	Driver
X-123197891-T-A	208	STAG2	STAG2 p.F672Y	Pair24	VUS
X-140969583-G-T	287	MAGEC3	MAGEC3 p.?	Pair24	VUS
1-36932452-G-C	227	CSF3R	CSF3R p.L700V	Pair25	VUS
1-51439893-T-A	394	CDKN2C	CDKN2C p.V153D	Pair25	VUS
11-119146845-G-A	328	CBL	CBL p.?	Pair25	Driver
11-119156151-A-G	221	CBL	CBL p.S606G	Pair25	Driver
11-62364125-T-A	302	MTA2	MTA2 p.D289V	Pair25	VUS
12-111856326-C-T	251	SH2B3	SH2B3 p.P126L	Pair25	VUS
12-111885286-C-T	56	SH2B3	SH2B3 p.R392W	Pair25	Driver
15-42046674-G-A	213	MGA	MGA p.V2350M	Pair25	VUS
16-30745085-C-T	134	SRCAP	SRCAP p.R2154*	Pair25	Driver
16-3786782-T-C	249	CREBBP	CREBBP p.S1477G	Pair25	VUS
16-57394357-G-A	5	CCL22	CCL22 p.G28S	Pair25	VUS
16-67650758-A-G	340	CTCF	CTCF p.M355V	Pair25	VUS

17-29665144-G-A	146	NF1	NF1 p.R2269H	Pair25	VUS
17-29670138-G-A	205	NF1	NF1 p.G2392R	Pair25	VUS
17-40477009-C-G	197	STAT3	STAT3 p.W479S	Pair25	VUS
2-25463295-T-G	262	DNMT3A	DNMT3A p.E733A	Pair25	Driver
2-25469633-G-A	140	DNMT3A	DNMT3A p.R379C	Pair25	Driver
2-25470021-C-T	96	DNMT3A	DNMT3A p.V341I	Pair25	Driver
2-65571975-C-T	162	SPRED2	SPRED2 p.G28R	Pair25	VUS
22-41527481-C-T	335	EP300	EP300 p.Q458*	Pair25	Possible
22-41573825-G-A	197	EP300	EP300 p.G2037D	Pair25	VUS
3-41278186-A-G	59	CTNNB1	CTNNB1 p.M688V	Pair25	VUS
3-47127765-G-A	271	SETD2	SETD2 p.R1773C	Pair25	VUS
3-47162156-G-A	363	SETD2	SETD2 p.Q1324*	Pair25	Driver
4-106155386-G-A	522	TET2	TET2 p.R96H	Pair25	Driver
4-153332496-C-T	57	FBXW7	FBXW7 p.V154I	Pair25	VUS
6-401645-G-A	266	IRF4	IRF4 p.G323R	Pair25	VUS
9-139404329-C-T	225	NOTCH1	NOTCH1 p.C942Y	Pair25	VUS
1-27087485-C-T	201	ARID1A	ARID1A p.P687S	Pair28	VUS
12-46245036-C-T	446	ARID2	ARID2 p.Q1044*	Pair28	Driver
2-25457192-G-C	249	DNMT3A	DNMT3A p.R899G	Pair28	Driver
21-36421183-C-T	237	RUNX1	RUNX1 p.S5N	Pair28	VUS
22-29083956-G-A	429	CHEK2	CHEK2 p.R564W	Pair28	Driver
22-29090060-C-T	414	CHEK2	CHEK2 p.R517H	Pair28	Driver
3-41266573-C-A	202	CTNNB1	CTNNB1 p.R124S	Pair28	VUS
3-47147564-C-T	201	SETD2	SETD2 p.E1588K	Pair28	VUS
3-47162848-C-T	28	SETD2	SETD2 p.S1093N	Pair28	VUS
4-106190818-C-T	215	TET2	TET2 p.R1366C	Pair28	Driver
4-106158085-GC-G	252	TET2	TET2 p.A996fs*11	Pair28	Driver
X-154348284-GA-G	261	BRCC3	BRCC3 p.N271fs*45	Pair28	Driver
1-120497790-T-C	131	NOTCH2	NOTCH2 p.I698V	Pair31	VUS
1-120512209-C-T	125	NOTCH2	NOTCH2 p.A345T	Pair31	VUS
11-118374730-A-C	119	KMT2A	KMT2A p.E2708A	Pair31	VUS
14-95574698-C-T	8	DICER1	DICER1 p.R800K	Pair31	VUS
16-30735573-C-T	146	SRCAP	SRCAP p.P1610S	Pair31	VUS
16-3823897-G-A	175	CREBBP	CREBBP p.P773L	Pair31	VUS
17-29548893-A-G	151	NF1	NF1 p.D556G	Pair31	VUS
17-58740550-G-T	27	PPM1D	PPM1D p.R485S	Pair31	Driver
19-44662159-C-T	60	ZNF234	ZNF234 p.R664*	Pair31	Driver
2-25505420-C-T	41	DNMT3A	DNMT3A p.G113E	Pair31	Driver
2-25994403-G-A	25	ASXL2	ASXL2 p.S137L	Pair31	VUS
X-39932819-C-T	86	BCOR	BCOR p.V594I	Pair31	Driver
22-30061020-GT-G	13	NF2	NF2 p.F285fs*11	Pair31	Driver
15-42032261-A-G	106	MGA	MGA p.N1482S	Pair38	VUS
17-7577124-C-T	366	TP53	TP53 p.V272M	Pair38	Driver
2-25468888-C-A	394	DNMT3A	DNMT3A p.?	Pair38	Driver
2-25470484-C-T	300	DNMT3A	DNMT3A p.W330*	Pair38	Driver
2-25470516-G-A	272	DNMT3A	DNMT3A p.R320*	Pair38	Driver
2-29455283-A-G	201	ALK	ALK p.I840T	Pair38	VUS
20-31390276-G-A	443	DNMT3B	DNMT3B p.R744K	Pair38	VUS
20-31390276-G-C	163	DNMT3B	DNMT3B p.R744T	Pair38	VUS
3-105470368-C-A	167	CBLB	CBLB p.D221Y	Pair38	VUS

4-106180807-G-T	149	<i>TET2</i>	TET2 p.E1279*	Pair38	Driver
6-43316334-T-C	206	<i>ZNF318</i>	ZNF318 p.K934E	Pair38	Driver
X-39933205-A-G	42	<i>BCOR</i>	BCOR p.V465A	Pair38	Driver
1-36933786-A-G	457	<i>CSF3R</i>	CSF3R p.I538T	Pair40	VUS
11-119145612-C-T	130	<i>CBL</i>	CBL p.T273M	Pair40	Driver
11-85975280-G-T	115	<i>EED</i>	EED p.G234V	Pair40	VUS
16-30727487-A-G	57	<i>SRCAP</i>	SRCAP p.Q865R	Pair40	VUS
17-29663364-T-C	16	<i>NF1</i>	NF1 p.L2007P	Pair40	VUS
17-7577566-T-C	36	<i>TP53</i>	TP53 p.N239D	Pair40	Driver
2-25457231-G-C	253	<i>DNMT3A</i>	DNMT3A p.Q886E	Pair40	Driver
2-65571995-A-T	26	<i>SPRED2</i>	SPRED2 p.M21K	Pair40	VUS
20-57428516-G-A	278	<i>GNAS</i>	GNAS p.E66K	Pair40	VUS
22-29091842-G-A	72	<i>CHEK2</i>	CHEK2 p.S415F	Pair40	Driver
22-29099506-T-C	355	<i>CHEK2</i>	CHEK2 p.I342V	Pair40	Driver
5-149460421-G-T	88	<i>CSF1R</i>	CSF1R p.N72K	Pair40	VUS
6-43305603-C-T	125	<i>ZNF318</i>	ZNF318 p.E2045K	Pair40	VUS
6-43333180-T-C	126	<i>ZNF318</i>	ZNF318 p.?	Pair40	Driver
7-148524346-C-T	69	<i>EZH2</i>	EZH2 p.R213H	Pair40	VUS
X-76938554-C-A	48	<i>ATRX</i>	ATRX p.D732Y	Pair40	VUS
4-106156053-AC-A	202	<i>TET2</i>	TET2 p.P319fs*28	Pair40	Driver
13-48951138-G-A	109	<i>RB1</i>	RB1 p.V434M	Pair41	VUS
16-30718975-G-A	259	<i>SRCAP</i>	SRCAP p.R192H	Pair41	Driver
2-26068405-T-C	18	<i>ASXL2</i>	ASXL2 p.M29V	Pair41	VUS
4-106197336-A-G	226	<i>TET2</i>	TET2 p.N1890S	Pair41	Driver
4-1919899-C-T	198	<i>WHSC1</i>	WHSC1 p.A320V	Pair41	VUS
4-1980493-G-A	26	<i>WHSC1</i>	WHSC1 p.G1319R	Pair41	VUS
5-131931498-A-G	195	<i>RAD50</i>	RAD50 p.M735V	Pair41	VUS
9-37020654-C-A	162	<i>PAX5</i>	PAX5 p.C64F	Pair41	VUS
X-53423265-A-G	64	<i>SMC1A</i>	SMC1A p.I915T	Pair41	VUS
6-43310585-AG-A	175	<i>ZNF318</i>	ZNF318 p.T1035fs*80	Pair41	Driver