

Supplemental information

**Direct cardiac reprogramming via combined
CRISPRa-mediated endogenous Gata4 activation
and exogenous Mef2c and Tbx5 expression**

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Table S1. Sequence, location and activation efficiency of Gata4 sgRNA in MEF and FCF

	sgRNA sequence	Location	Gata4 endo in MEF			Gata4 total in MEF			Gata4 endo in FCF			Gata4 total in FCF		
G1	GGATCTCAGGAAAATCCCAG	chr14:63246528	2.14	±	0.30	1.69	±	0.03	1.94	±	0.07	1.75	±	0.20
G2	GGAGTGGGAAGAAGTGTGG	chr14:63245877	129.21	±	33.39	174.25	±	8.74	1.40	±	0.10	1.65	±	0.10
G3	GTGGACCAC TGAGAGTAGGG	chr14:63245551	132.10	±	15.62	191.68	±	20.51	1.49	±	0.19	1.76	±	0.40
G4	GTCGGTCAGAGAGATTACAC	chr14:63244811	2.14	±	0.30	1.69	±	0.03	0.10	±	0.02	0.09	±	0.02
G5	TTGCTAGCCTCAGATCTACG	chr14:63245674	1.94	±	0.07	1.75	±	0.20	1.40	±	0.10	1.65	±	0.10
G6	GGGCTGC ACTGAGGGCAGAA	chr14:63245473	226.72	±	18.97	302.77	±	33.00	0.90	±	0.08	2.50	±	0.09
G7	GGGGCGGGGGAGCCCGGACC	chr14:63245339	31.58	±	2.48	43.15	±	9.42	1.44	±	0.15	1.73	±	0.04
G8	GCCTAAGGGAGTCACGTGCA	chr14:63245313	3.16	±	0.33	2.12	±	0.39	0.93	±	0.11	0.78	±	0.04
G9	GGGGCCCGGGGAACCGCGCC	chr14:63245264	37.30	±	2.55	30.64	±	2.59	1.14	±	0.15	1.32	±	0.09
G10	CGCCGGCGGGAGGTGCTGCC	chr14:63245357	284.69	±	18.10	311.04	±	5.91	1.82	±	0.13	1.72	±	0.04
G11	GCATGGACTTGCCTGTTGG	chr14:63245407	0.93	±	0.05	1.21	±	0.05	0.10	±	0.02	0.09	±	0.02
G12	TTGGGAAGAGTCCTGCGGGC	chr14:63245380	23.25	±	1.90	30.66	±	1.15	1.29	±	0.07	1.57	±	0.13
G13	AGCGCAGGCGATCGCTACGC	chr14:63245285	20.13	±	1.46	25.72	±	1.40	1.44	±	0.15	1.73	±	0.04
G14	GAGTCCTGCAGGGCGGGCGCC	chr14:63245373	14.07	±	2.19	14.92	±	1.20	0.82	±	0.04	0.89	±	0.10
G15	CGCAGGCGATCGCTACGCGG	chr14:63245283	20.81	±	2.20	14.04	±	2.57	0.95	±	0.17	0.81	±	0.05

Table S2. Sequence, location and activation efficiency of Mef2c sgRNA in MEF and FCF

	sgRNA sequence	Location	Mef2c endo in MEF			Mef2c total in MEF		Mef2c endo in FCF		Mef2c total in FCF				
M1	AAATGAGCTCGGGCAAAGAA	chr13:83503732	0.62	±	0.05	0.65	±	0.04	0.48	±	0.02	0.46	±	0.03
M2	AAGAGACTCGGTGTCAAAAC	chr13:83503618	0.60	±	0.07	0.55	±	0.06	0.44	±	0.01	0.35	±	0.02
M4	TCCAATGGAAAATATCCAAT	chr13:83503980	0.51	±	0.05	0.40	±	0.02	0.64	±	0.05	0.46	±	0.05
M5	TACATTCCAAGAATAATCT	chr13:83503868	0.39	±	0.04	0.40	±	0.06	0.42	±	0.01	0.31	±	0.09
M6	GTGGCTGGAAACTTTTAA	chr13:83503818	0.56	±	0.04	0.50	±	0.03	0.70	±	0.03	0.55	±	0.05
M7	GAAAAAAAGCAAATGAGCTG	chr13:83503722	0.68	±	0.01	0.61	±	0.21	1.03	±	0.05	0.95	±	0.25
M8	GCTACTGTACCATTAAAA	chr13:83503560	0.42	±	0.00	0.39	±	0.03	0.57	±	0.05	0.41	±	0.06
M9	CCGATTGGATATTTCCAT	chr13:83503979	0.53	±	0.03	0.53	±	0.06	4.68	±	0.22	2.56	±	0.12
M10	AGTTACAAGCTTCTAATT	chr13:83503643	0.75	±	0.00	0.55	±	0.03	0.62	±	0.03	0.46	±	0.03
M11	TGAAAAAAAGCAAATGAGCTG	chr13:83503722	0.85	±	0.11	0.89	±	0.13	0.66	±	0.00	0.71	±	0.04
M12	AAAATGGACACTCACGTCTG	chr13:83503544	0.98	±	0.05	0.96	±	0.14	0.80	±	0.10	0.78	±	0.08
M13	GACTCGGTGTCAAAACTGGA	chr13:83503614	1.81	±	0.09	1.88	±	0.16	0.70	±	0.01	0.76	±	0.09
M14	ACTAACAGTGTAGAGGCTTG	chr13:83504003	0.97	±	0.32	1.05	±	0.31	0.68	±	0.04	0.79	±	0.03
M15	TAAATGGTACAGTAGCATTG	chr13:83503566	0.80	±	0.02	0.85	±	0.10	0.84	±	0.11	0.91	±	0.18
M16	GTTAAAACCTTATCACATAGG	chr13:83503679	1.27	±	0.09	1.36	±	0.10	0.98	±	0.04	0.95	±	0.06
M17	TTTAAAGCCTGTGTGAAATG	chr13:83503893	0.59	±	0.03	0.64	±	0.11	1.07	±	0.01	1.00	±	0.21
M18	AGTTTTGTTCAATTAAATTG	chr13:83503836	1.12	±	0.12	1.12	±	0.23	0.75	±	0.05	0.88	±	0.07
M21	GGTCATGGCACTTAAACGAT	chr13:83503593	0.69	±	0.01	0.71	±	0.04	0.89	±	0.10	1.00	±	0.10

M22	CAATCCAAAGAAATTAGA	chr13:83503701	0.77	\pm	0.06	0.79	\pm	0.08	0.84	\pm	0.03	0.99	\pm	0.25
M23	GTTGGCTTCAGTCTGGTCG	chr13:83502595	0.38	\pm	0.05	0.34	\pm	0.05	0.58	\pm	0.04	0.65	\pm	0.01
M24	TACATAAGAACATGAGACCTGA	chr13:83502807	0.44	\pm	0.02	0.37	\pm	0.05	0.35	\pm	0.05	0.36	\pm	0.05
M25	AGAGAAAGTACACAGTGAGAG	chr13:83503012	0.81	\pm	0.04	0.72	\pm	0.06	0.41	\pm	0.06	0.41	\pm	0.02
M26	GTCTATGTTTATCTGAAAG	chr13:83503076	0.43	\pm	0.14	0.40	\pm	0.12	0.50	\pm	0.09	0.64	\pm	0.17
M27	TTAAAACGTGAGAAAAGTTA	chr13:83502652	0.35	\pm	0.01	0.33	\pm	0.04	0.30	\pm	0.00	0.54	\pm	0.06
M28	AAGAACATGAGACCTGATGGAG	chr13:83502812	0.56	\pm	0.04	0.52	\pm	0.04	0.31	\pm	0.06	0.38	\pm	0.07
M29	TATTTAATAATTATACGGGT	chr13:83502942	0.26	\pm	0.01	0.25	\pm	0.04	0.41	\pm	0.08	0.42	\pm	0.10
M30	CAGTGAGAGAGGTGCTTGCA	chr13:83503023	0.50	\pm	0.05	0.43	\pm	0.09	0.58	\pm	0.17	0.46	\pm	0.10
M31	GACACAAGGCCTTGAAAAG	chr13:83502714	0.31	\pm	0.00	0.27	\pm	0.02	0.22	\pm	0.01	0.19	\pm	0.04
M32	ATGAGACCTGATGGAGAGGT	chr13:83502816	0.34	\pm	0.03	0.30	\pm	0.03	0.45	\pm	0.04	0.35	\pm	0.11
M33	GGAGCTTGCTAAAAAGAAC	chr13:83502911	0.29	\pm	0.00	0.27	\pm	0.02	0.31	\pm	0.03	0.29	\pm	0.05
M34	ATGGCATCACCCAGTATCCA	chr13:83503049	0.36	\pm	0.04	0.30	\pm	0.03	0.20	\pm	0.02	0.18	\pm	0.03
M35	TTTGTTAGGCCACTTTCAA	chr13:83502723	0.31	\pm	0.01	0.29	\pm	0.04	0.29	\pm	0.02	0.28	\pm	0.04
M36	ATGAAATATGAGGAAACCTA	chr13:83502857	0.30	\pm	0.02	0.30	\pm	0.01	0.30	\pm	0.06	0.32	\pm	0.08
M37	AATGGCATCACCCAGTATCC	chr13:83503050	0.38	\pm	0.05	0.35	\pm	0.07	0.44	\pm	0.02	0.39	\pm	0.16
M38	TCTGAAAGTGGAGCCCTGCA	chr13:83503088	0.43	\pm	0.02	0.36	\pm	0.02	0.29	\pm	0.01	0.31	\pm	0.04
M39	AAAAGTGGCCTAACAAATAT	chr13:83502729	0.47	\pm	0.00	0.38	\pm	0.08	0.29	\pm	0.01	0.28	\pm	0.01
M40	TGAAATATGAGGAAACCTAA	chr13:83502858	0.33	\pm	0.02	0.34	\pm	0.03	0.26	\pm	0.02	0.22	\pm	0.07
M41	GATACTGGGTGATGCCATT	chr13:83503054	0.39	\pm	0.05	0.38	\pm	0.04	0.40	\pm	0.09	0.34	\pm	0.09
M42	TGGAGCCCTGCAGGGAAATG	chr13:83503096	0.37	\pm	0.02	0.38	\pm	0.10	0.18	\pm	0.02	0.15	\pm	0.02

M43	ATCACACACCAATATTGTT	chr13:83502737	0.26	±	0.02	0.25	±	0.00	0.32	±	0.09	0.36	±	0.09
M44	TGCGTCATAACAAAACCCTT	chr13:83502873	0.23	±	0.01	0.18	±	0.03	0.81	±	0.19	0.96	±	0.24
M45	TTATTTAATAATTATACGGG	chr13:83502943	0.27	±	0.03	0.24	±	0.01	0.31	±	0.11	3.57	±	0.18
M46	AGATAAAACATAGACCTGAA	chr13:83503068	0.20	±	0.00	0.16	±	0.02	0.62	±	0.09	1.88	±	0.49
M47	ACAAGCCTCATTCCCTGCA	chr13:83503101	0.52	±	0.02	0.47	±	0.07	0.60	±	0.06	0.61	±	0.04
M49	TCTTGTCCAAGATTATTCT	chr13:83503861	0.85	±	0.17	0.64	±	0.10	1.88	±	0.25	1.73	±	0.11
M50	CAGTGTAGAGGCTGGGTG	chr13:83504008	0.87	±	0.19	0.84	±	0.08	1.40	±	0.10	1.14	±	0.23
M51	TCTTTGCCAGCACTGACAA	chr13:83504048	0.59	±	0.05	0.48	±	0.05	0.16	±	0.02	0.19	±	0.01
M52	CGGTGTTCATAGAAAAGGAG	chr13:83504202	0.74	±	0.04	0.64	±	0.11	0.85	±	0.06	0.58	±	0.02
M53	TGCACCTGTTCATGTCACTC	chr13:83504108	0.50	±	0.08	5.04	±	0.36	0.23	±	0.04	0.22	±	0.03
M54	AATAGCACATGGAATTTTG	chr13:83504798	0.86	±	0.14	0.77	±	0.14	0.77	±	0.02	0.54	±	0.01
M55	TGCCAGCACTGACAAAGGTC	chr13:83504053	0.54	±	0.04	0.57	±	0.10	0.21	±	0.01	0.18	±	0.03
M56	TACTCCAGAGTGACATGAAC	chr13:83504104	1.18	±	0.03	1.03	±	0.02	1.29	±	0.05	1.25	±	0.09
M57	GTTCCCGTCAGCACCTGCTG	chr13:83504326	0.55	±	0.01	0.57	±	0.10	0.30	±	0.01	0.29	±	0.01
M58	AAGGAAGCAGCTCAAAGCTA	chr13:83505263	1.05	±	0.14	1.00	±	0.14	1.17	±	0.12	1.07	±	0.05
M60	GGCGAGCGCAGCCAACTG	chr13:83504226	0.74	±	0.16	0.83	±	0.11	0.13	±	0.00	0.21	±	0.02
M61	TGCTGACGGGAACAACTTCC	chr13:83504336	0.40	±	0.05	0.52	±	0.06	0.23	±	0.02	0.22	±	0.02
M62	ATGGAATTTACTTATTAAAA	chr13:83505310	0.79	±	0.10	0.43	±	0.04	1.12	±	0.07	1.07	±	0.09
M64	TGAAGAGAAACCCCCCAGTT	chr13:83504238	0.68	±	0.05	0.63	±	0.08	0.24	±	0.02	0.19	±	0.03
M65	ATCATCAGCCTTGTAACAG	chr13:83504463	0.61	±	0.07	0.56	±	0.05	0.56	±	0.04	0.52	±	0.03
M66	TTCCATATTACAAGCAACA	chr13:83505327	0.66	±	0.01	0.63	±	0.15	0.23	±	0.02	0.22	±	0.02

M67	GCACAAGTGTCTGGCAGGCC	chr13:83504132	0.58	±	0.02	0.53	±	0.10	0.19	±	0.01	0.17	±	0.01
M68	TAGAATAAAGCCAGACCAGC	chr13:83504264	0.42	±	0.03	0.56	±	0.05	0.23	±	0.01	0.27	±	0.01
M69	AAGGCTGATGATGAGTGAGC	chr13:83504474	0.45	±	0.05	0.52	±	0.05	0.23	±	0.01	0.27	±	0.01
M70	AATGAATGTAAAAGACACAA	chr13:83505353	0.55	±	0.02	0.46	±	0.07	0.77	±	0.03	0.79	±	0.03
M71	CTGCCAGACACTTGTGCAGA	chr13:83504138	0.48	±	0.02	0.39	±	0.06	0.68	±	0.11	0.75	±	0.03
M72	GAGGCGAGCGCAGCCCCAAC	chr13:83504224	0.28	±	0.01	0.49	±	0.04	0.20	±	0.01	0.19	±	0.02
M73	GCACCTAGCAACCCCACGT	chr13:83504514	0.71	±	0.07	0.51	±	0.11	0.31	±	0.00	0.34	±	0.03
M74	TCAATTCTGGGCTGTGGG	chr13:83505418	0.88	±	0.02	0.23	±	0.01	0.88	±	0.02	0.78	±	0.07
M76	CGAGCGCAGCCAACTGGG	chr13:83504228	0.94	±	0.15	0.60	±	0.13	0.23	±	0.02	0.22	±	0.01
M77	TTCTTAGAGGAGACAGTGC	chr13:83504436	0.82	±	0.04	0.78	±	0.07	0.30	±	0.03	0.30	±	0.04
M78	TGGGCTGTGGGGGTAGGAT	chr13:83505427	0.62	±	0.05	0.93	±	0.08	0.63	±	0.07	0.59	±	0.08
M80	TTTGTGTCACAACTCTGAC	chr13:83504301	0.59	±	0.08	0.84	±	0.14	0.21	±	0.04	0.20	±	0.03
M81	GATGATGAGTGAGCTGGAAA	chr13:83504480	0.60	±	0.02	0.78	±	0.07	0.28	±	0.01	0.26	±	0.01
M82	TTACTGTTATTATAAAGGA	chr13:83505820	0.60	±	0.07	0.58	±	0.07	0.60	±	0.07	0.52	±	0.03
M84	TGGACAACAAAGCCCTCAGC	chr13:83504313	0.77	±	0.03	0.67	±	0.07	0.22	±	0.00	0.22	±	0.03
M85	GGAGCCAACCTCCCCAACAG	chr13:83504501	0.96	±	0.12	0.55	±	0.07	0.29	±	0.01	0.29	±	0.03
M86	CTTTACTTAAGGCTTCGCAT	chr13:83505874	1.50	±	0.03	0.56	±	0.05	1.50	±	0.03	1.32	±	0.18
M88	AAGCCCTCAGCAGGTGCTGA	chr13:83504322	0.79	±	0.03	0.52	±	0.03	0.23	±	0.01	0.21	±	0.01
M89	ACAAACGCCAACACATTGAA	chr13:83504542	0.74	±	0.01	0.72	±	0.06	0.72	±	0.10	0.53	±	0.05
M90	ACTTATTGCTGATTCTGAGA	chr13:83505930	0.81	±	0.04	0.72	±	0.06	0.81	±	0.04	0.78	±	0.07
M91	TGGAGTAAATTAGCTGTAA	chr13:83504088	0.46	±	0.07	0.99	±	0.20	0.46	±	0.07	0.53	±	0.05

M92	CTCACCGCTTGACGATCAAG	chr13:83504170	ME	0.46	±	0.06	0.42	±	0.04	1.02	±	0.03	1.09	±	0.06
M93	GATCCCTCTGCACAAGTGTG	chr13:83504141	ME	0.58	±	0.02	0.53	±	0.06	1.12	±	0.03	1.12	±	0.17
M94	CTCCCCCTGACCGATAGATAG	chr13:83523356	HFE	1.41	±	0.07	1.27	±	0.08	2.48	±	0.16	91.44	±	12.36
M95	AGTAAGGTGTGGAGGGAAAGG	chr13:83523436	HFE	0.66	±	0.08	0.56	±	0.06	1.40	±	0.05	1.51	±	0.07
M96	TCACCGCTTGACGATCAAGG	chr13:83504171	ME	0.62	±	0.02	0.52	±	0.04	1.01	±	0.03	1.12	±	0.06
M97	TAAGTTTCCTCATTCACAC	chr13:83503900	ME	0.74	±	0.02	0.64	±	0.05	0.80	±	0.07	0.78	±	0.06
M98	TGGTTTACTTGCTAATGACC	chr13:83523404	HFE	0.38	±	0.01	0.34	±	0.01	1.20	±	0.00	1.25	±	0.08
M99	TACACTTGTGGAGCAGTTA	chr13:83572135	ESE	0.46	±	0.06	0.43	±	0.03	1.30	±	0.09	1.35	±	0.08
M100	ATTTTGGATAGACTTCCGAT	chr13:83503965	ME	0.48	±	0.02	0.41	±	0.05	1.08	±	0.02	1.16	±	0.11
M101	CATTGGAACTAACAGTGTAG	chr13:83503996	ME	0.92	±	0.05	0.85	±	0.10	0.83	±	0.02	0.86	±	0.04
M102	TCAGGGGAGCCTAACATGCATT	chr13:83523370	HFE	1.60	±	0.09	1.43	±	0.31	2.99	±	0.06	3.32	±	0.20
M103	CAGCAACCGCGAACAAATAAA	chr13:83572175	ESE	0.50	±	0.04	0.46	±	0.04	1.11	±	0.03	1.12	±	0.02
M104	TTGCCCCCTTGATCGTCAAG	chr13:83504174	ME	0.49	±	0.01	0.43	±	0.05	1.23	±	0.18	38.58	±	9.25
M105	TGTGGCTGGAAACTTTTAA	chr13:83503818	ME	0.51	±	0.03	0.47	±	0.05	1.05	±	0.03	1.03	±	0.09
M106	GAAATCTGACTTTATCCCC	chr13:83523522	HFE	0.35	±	0.04	0.34	±	0.07	1.21	±	0.08	1.29	±	0.08
M107	GAAACACGCACAGACTGGCC	chr13:83572212	ESE	0.51	±	0.01	0.42	±	0.01	0.91	±	0.01	0.94	±	0.05
M108	CGATCAAGGGGGCAAAGCTT	chr13:83504182	ME	0.66	±	0.06	0.63	±	0.03	1.03	±	0.02	1.05	±	0.07
M109	ACCCGCTATCTATCGGTAG	chr13:83523354	HFE	0.87	±	0.09	0.77	±	0.02	1.85	±	0.02	2.14	±	0.10
M110	ATCTGGACAAATTACTGAG	chr13:83523481	HFE	0.61	±	0.09	0.50	±	0.08	1.20	±	0.02	1.23	±	0.08
M111	AGATTCTCTGTTACTAGGA	chr13:83572249	ESE	0.54	±	0.05	0.49	±	0.02	1.33	±	0.05	1.27	±	0.04
M112	AACCAGACCTTGTCA GTGC	chr13:83504055	ME	0.60	±	0.08	0.50	±	0.01	0.93	±	0.02	1.15	±	0.11

M113	CACCCGCTATCTATCGGTCA	chr13:83523353	HFE	1.36	\pm	0.15	1.18	\pm	0.17	2.65	\pm	0.15	2.75	\pm	0.04
M114	GGATAAAGTCCAGTAAGGTG	chr13:83523425	HFE	0.64	\pm	0.07	0.62	\pm	0.05	1.20	\pm	0.02	1.30	\pm	0.12
M115	ACTGCTCCACAAGTGTAAAA	chr13:83572129	ESE	0.52	\pm	0.04	0.46	\pm	0.02	0.98	\pm	0.03	1.02	\pm	0.04
M116	AGCTTCGGTGTTCATAGAAA	chr13:83504197	ME	0.48	\pm	0.05	0.46	\pm	0.03	0.94	\pm	0.11	0.78	\pm	0.10
M117	ATTGAAAGTTAACGGCCCG	chr13:83523506	HFE	0.78	\pm	0.08	0.71	\pm	0.13	0.86	\pm	0.05	0.73	\pm	0.09
M118	GACCTGGATAAAGTCCAGTA	chr13:83523420	HFE	0.49	\pm	0.01	0.46	\pm	0.06	0.72	\pm	0.04	0.60	\pm	0.04
M120	TGACATGAACAGGTGCACCC	chr13:83504114	ME	1.34	\pm	0.03	1.25	\pm	0.06	0.85	\pm	0.07	7.47	\pm	1.06
M121	TGAAGTCACCCGCTATCTAT	chr13:83523347	HFE	0.68	\pm	0.09	0.63	\pm	0.03	1.68	\pm	0.04	1.31	\pm	0.16
M122	GTTCAGCTGCACAGGAAGC	chr13:83523311	HFE	0.23	\pm	0.01	0.22	\pm	0.01	1.08	\pm	0.06	0.90	\pm	0.08
M123	ACGCACAGACTGCCAGGGA	chr13:83572207	ESE	0.56	\pm	0.00	0.54	\pm	0.02	0.79	\pm	0.03	0.71	\pm	0.04
M124	GAGAAGGAAGTGGAGAGTTT	chr13:83572103	ESE	1.24	\pm	0.02	1.15	\pm	0.09	1.00	\pm	0.07	1.08	\pm	0.13

ME, Muscle enhancer; HFE, Heart field enhancer; ESE, Endothelial cell-specific enhancer

Table S3. Sequence, location and activation efficiency of Tbx5 sgRNA in MEF and FCF

	sgRNA sequence	Location	Tbx5 endo in MEF		Tbx5 total in MEF		Tbx5 endo in FCF		Tbx5 total in FCF					
T1	CTGAGCCAGTGGTTGCAGGG	chr5:119834446	2.58	±	1.10	2.45	±	1.12	0.35	±	0.03	0.71	±	0.04
T2	AAGTCCTAGAGAGCTTGGAG	chr5:119834094	3.05	±	0.77	2.31	±	0.29	3.67	±	2.95	1.06	±	0.09
T5	GGACAATGAGTCTGAAGTGG	chr5:119834128	5.56	±	1.75	4.07	±	0.86	37.07	±	1.65	11.22	±	1.41
T6	GAAATCGGGTGAGGCTGCAG	chr5:119834236	1.93	±	0.42	0.71	±	0.03	7.04	±	2.15	1.33	±	0.09
T7	AGGAAGGAAGGAAAGAAGGA	chr5:119834374	2.39	±	0.89	1.21	±	0.32	38.13	±	7.87	10.26	±	1.16
T9	TATAGTGGTTCAAGAGTTG	chr5:119834546	0.49	±	0.40	1.04	±	0.03	0.92	±	0.13	0.95	±	0.03
T10	GCCCTGCAGAGAACCAAGAC	chr5:119834289	1.01	±	0.22	1.59	±	0.16	6.04	±	0.56	2.29	±	0.32
T11	TAGGCGTGTGCACACACCCA	chr5:119834260	0.78	±	0.09	0.61	±	0.09	3.23	±	0.62	1.02	±	0.07
T12	GCTAGTCCTGGCTCTGCAAG	chr5:119834166	5.10	±	0.54	3.45	±	0.48	8.53	±	0.99	2.28	±	0.28
T13	AGTGGGGGTGGGAATCAGC	chr5:119834520	1.60	±	0.67	0.85	±	0.05	0.28	±	0.09	0.50	±	0.08
T14	AAGAGGATGGGAAGTGGAAA	chr5:119834497	1.11	±	0.36	0.86	±	0.02	8.17	±	8.83	0.70	±	0.04
T15	AAAGTGGAAAGGTGGTGGG	chr5:119834582	1.55	±	0.37	3.01	±	0.08	16.20	±	15.07	0.66	±	0.06
T16	AGTCAACAGAGATGGAAGGA	chr5:119834472	34.74	±	4.50	15.05	±	2.30	29.63	±	33.36	1.00	±	0.22
T17	GGGAAGTGGAAAGGGAGTGG	chr5:119834505	1.55	±	0.21	1.23	±	0.01	0.32	±	0.04	0.64	±	0.16
T18	ATCTAGTGAGAAAGTGGAA	chr5:119834572	0.72	±	0.05	2.17	±	0.37	2.30	±	0.44	0.99	±	0.20

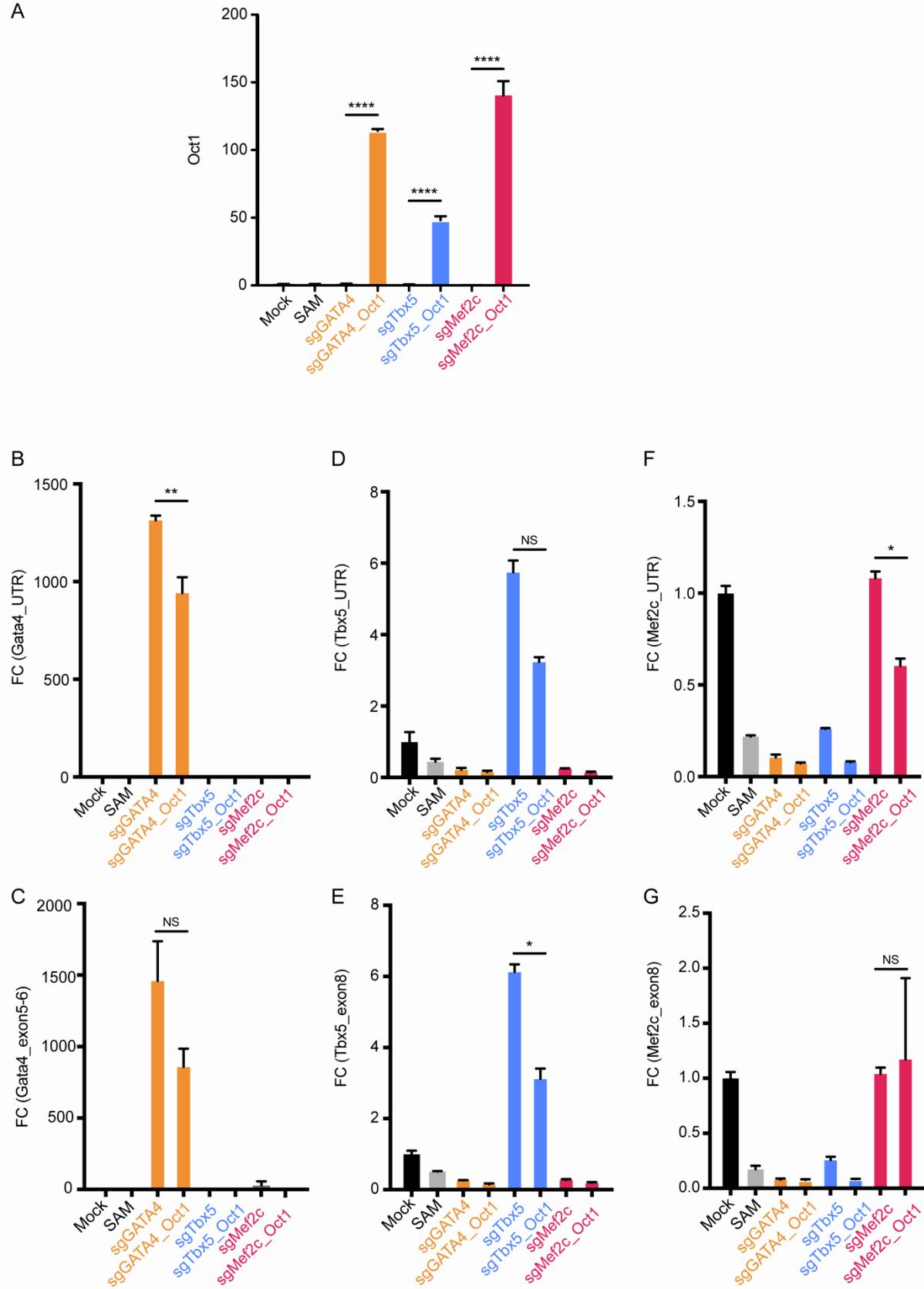


Figure S1. Gata4, Tbx5 and Mef2c activation level by adding Oct1 in MEF. (A) Transcriptional activation of Oct1 in MEFs with SAM system among mock, M/G/T sgRNA alone or Oct1 combination groups. (B) and (C) Transcriptional activation of endogenous Gata4 (B) and total Gata4 (C) with SAM activators. (D) and (E) Transcriptional activation of endogenous Tbx5 (D) and total Tbx5 (E) with SAM activators. (F) and (G) Transcriptional activation of endogenous Mef2c (F) and total Mef2c (G) with SAM activators.

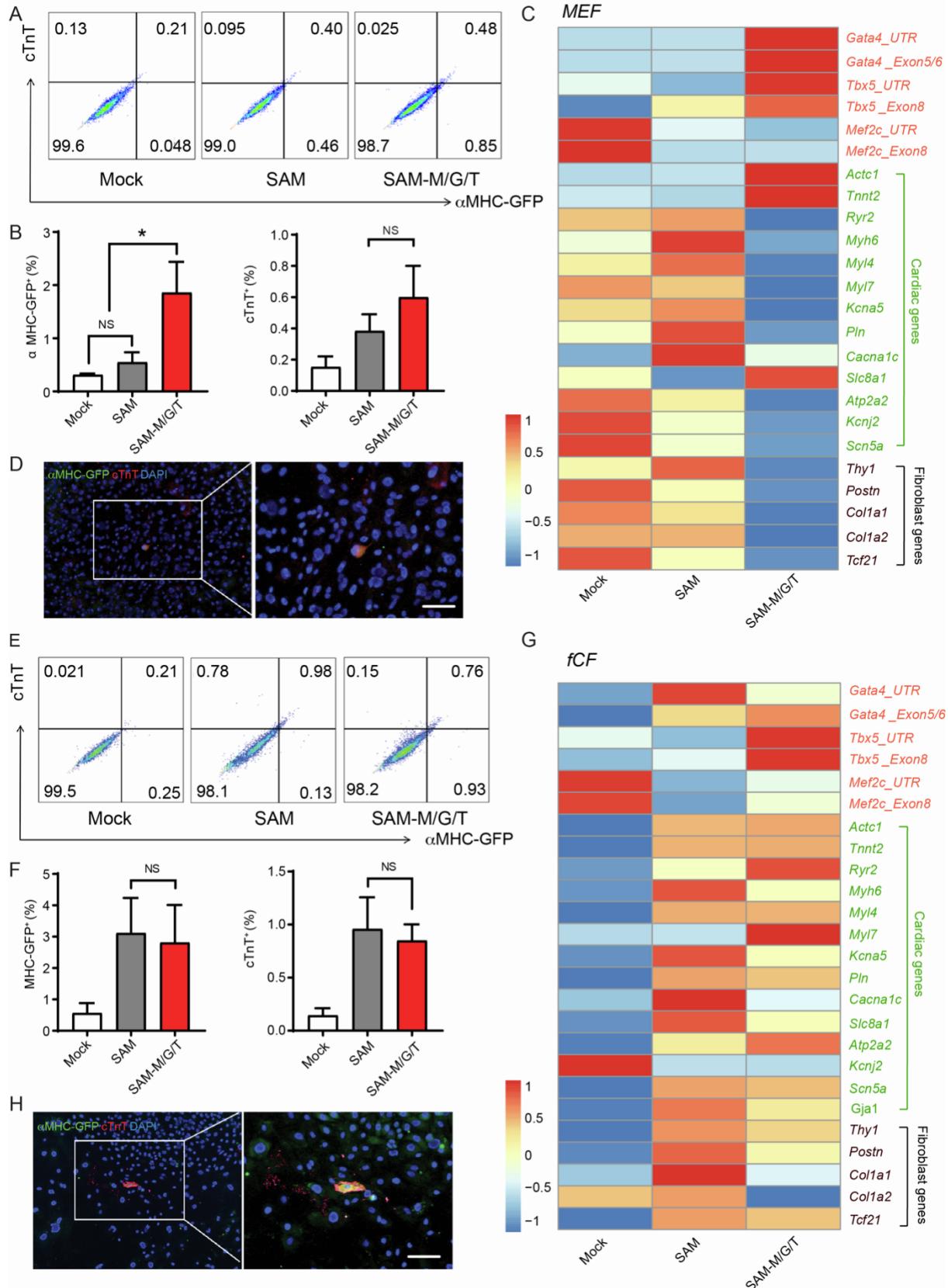


Figure S2. CRISPRa-mediated activation of endogenous *Mef2c*, *Gata4*, and *Tbx5* promotes cardiac gene expression and suppresses the fibroblast gene program in embryonic fibroblasts and fresh cardiac fibroblasts. (A) Flow cytometry analysis of GFP and cTnT in MEFs after reprogramming and the statistics graph of A (B). (C) Heat map of transcriptional activators, cardiac progenitor markers and fibrosis markers on day 10 post infection by M/G/T-sgRNA in MEFs. (D) Immunofluorescence labeling of GFP and cTnT on day 14 of reprogramming in MEFs. (E) Flow cytometry analysis of GFP and cTnT in fCFs after reprogramming and the statistics graph of E (F). (G) Heat map of transcriptional activators, cardiac progenitor markers and fibrosis markers on day 10 post infection by M/G/T-sgRNA in fCFs. (H) Immunofluorescence labeling of GFP and cTnT on day 14 of reprogramming in fCFs. Scale bars, 100 μ m (D and H). * $p < 0.05$.

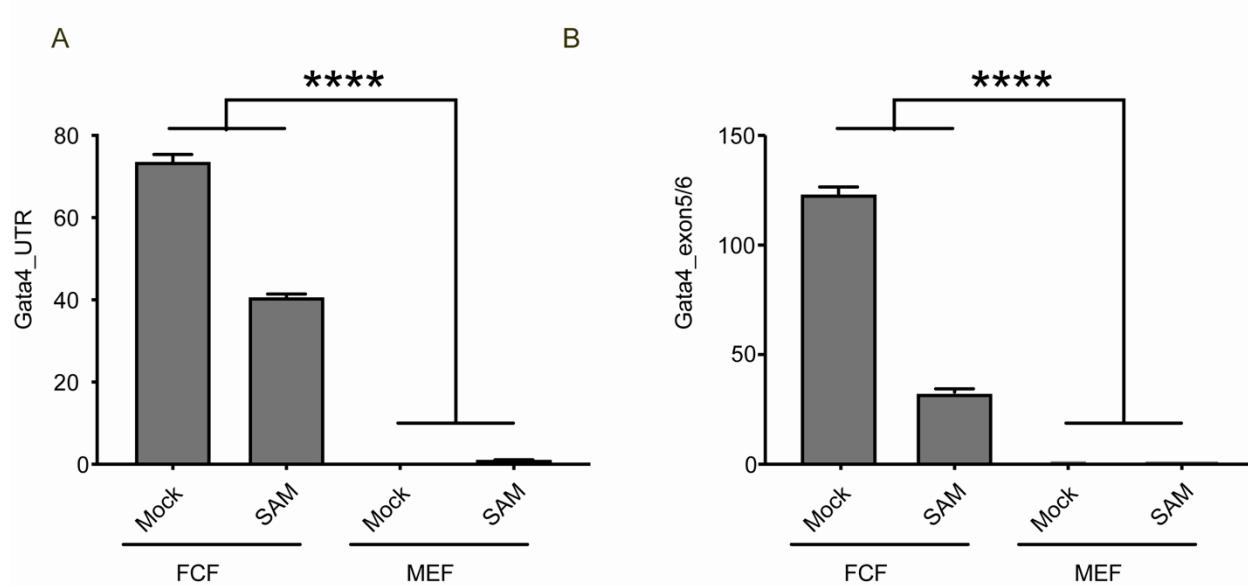


Figure S3. Basic expression level of endogenous and total Gata4 in fCFs and MEFs. (A) Expression level of endogenous Gata4 (Gata4_UTR) in fCFs vs. MEFs. (B) Expression level of total Gata4 (Gata4_exon5/6) in fCFs vs. MEFs. *** $p < 0.0001$.