

Gene Name	Description	Probe Set ID	Low-HbF						High-HbF					
			#1	#2	#3	#4	#5	Ave±SD	#1	#2	#3	#4	#5	Ave±SD
GATA1	GATA binding protein 1 (globin transcription factor 1)	1555590_a_at	16.1	9.2	25.3	17.5	15.9	16.8±5.7	17.6	12.9	15.4	10.5	10.9	13.5±3.0
	GATA binding protein 1 (globin transcription factor 1)	210446_at	20.0	19.3	19.0	19.0	22.2	19.9±1.3	19.0	17.7	19.1	18.3	15.5	17.9±1.5
GATA2	GATA binding protein 2	207954_at	2.4	0.6	0.9	1.5	2.0	1.5±0.7	1.9	1.0	2.3	1.2	1.9	1.7±0.6
	GATA binding protein 2	209710_at	2.2	2.6	3.2	2.2	3.3	2.7±0.5	4.4	5.0	5.9	4.8	5.0	5.0±0.6
	GATA binding protein 2	210358_x_at	2.4	0.6	3.0	2.3	3.0	2.3±1.0	3.4	2.7	2.5	4.2	3.0	3.1±0.7
GATA3	GATA binding protein 3	209602_s_at	0.1	0.1	0.0	0.1	0.0	0.1±0.0	0.0	0.0	0.0	0.1	0.1	0.1±0.0
	GATA binding protein 3	209603_at	0.1	0.2	0.1	0.2	0.1	0.1±0.0	0.1	0.2	0.5	0.1	0.1	0.2±0.2
	GATA binding protein 3	209604_s_at	2.0	0.1	0.6	0.2	0.8	0.7±0.8	1.2	0.6	1.1	0.8	0.2	0.8±0.4
GATA4	GATA binding protein 4	1553131_a_at	1.3	0.4	0.4	2.2	1.1	1.1±0.7	0.3	0.4	2.6	1.5	0.5	1.1±1.1
	GATA binding protein 4	1570276_a_at	0.1	0.1	0.9	0.1	0.3	0.3±0.4	0.2	0.2	0.5	0.6	0.2	0.3±0.2
	GATA binding protein 4	205517_at	0.5	0.6	0.1	0.1	0.6	0.4±0.2	0.4	0.5	0.1	0.1	0.2	0.3±0.2
	GATA binding protein 4	230855_at	0.2	1.1	1.5	0.3	0.3	0.7±0.6	1.4	2.2	0.2	1.0	0.1	1.0±0.9
	GATA binding protein 4	243692_at	0.5	1.6	1.0	1.5	0.7	1.0±0.4	1.3	1.0	0.8	1.1	0.8	1.0±0.2
GATA5	GATA binding protein 5	238095_at	0.4	1.6	0.6	1.5	0.7	1.0±0.5	2.8	0.2	1.6	2.5	1.6	1.7±1.0
	GATA binding protein 5	238197_at	1.1	1.3	1.0	1.8	0.7	1.2±0.4	3.2	2.4	2.6	2.7	2.5	2.7±0.3
GATA6	GATA binding protein 6	210002_at	0.1	0.7	0.1	1.0	0.7	0.5±0.4	0.0	0.1	0.5	0.1	0.3	0.2±0.2
	GATA binding protein 6	229282_at	2.2	0.9	1.0	1.5	1.5	1.4±0.5	1.2	1.3	1.4	0.2	1.4	1.1±0.5
KLF1 (EKLF)	Kruppel-like factor 1 (erythroid)	210504_at	77.7	69.4	96.7	75.4	90.3	81.9±11.2	63.6	55.7	68.5	58.0	38.9	56.9±11.2
KLF2 (LKLF)	Kruppel-like factor 2 (lung)	219371_s_at	0.5	1.0	0.1	0.2	0.2	0.4±0.4	4.4	3.3	3.7	1.0	4.2	3.3±1.4
	Kruppel-like factor 2 (lung)	226645_at	0.1	0.1	0.1	0.1	0.1	0.1±0.0	0.4	0.5	0.1	0.1	0.1	0.2±0.2
	Kruppel-like factor 2 (lung)	226646_at	0.2	0.7	0.8	1.5	1.3	0.9±0.5	1.5	1.2	0.5	0.6	0.3	0.8±0.5
KLF3 (BKLF)	Kruppel-like factor 3 (basic)	219657_s_at	6.8	4.9	5.6	5.2	7.6	6.0±1.1	4.3	6.5	3.5	4.6	5.8	5.0±1.2
	Kruppel-like factor 3 (basic)	222913_at	5.3	3.3	5.0	5.4	5.0	4.8±0.8	7.4	4.8	6.4	3.3	3.9	5.2±1.7
	Kruppel-like factor 3 (basic)	225133_at	66.6	45.1	51.3	54.4	64.8	56.4±9.1	44.7	38.8	42.2	34.1	37.0	39.4±4.2
	Kruppel-like factor 3 (basic)	225140_at	13.5	7.5	11.6	11.5	12.5	11.3±2.3	10.6	8.7	9.3	8.1	6.3	8.6±1.6
KLF4 (GKLF)	Kruppel-like factor 4 (gut)	220266_s_at	0.1	1.1	0.1	0.1	0.1	0.3±0.5	0.3	0.2	0.3	0.1	0.1	0.2±0.1
	Kruppel-like factor 4 (gut)	221841_s_at	1.1	1.7	1.2	1.0	1.0	1.2±0.3	3.8	4.0	3.6	3.5	2.5	3.5±0.6
KLF5 (IKLF)	Kruppel-like factor 5 (intestinal)	209211_at	0.8	0.6	1.0	0.6	1.3	0.9±0.3	0.5	0.3	2.3	0.0	0.4	0.7±0.9
	Kruppel-like factor 5 (intestinal)	209212_s_at	1.6	1.1	2.8	1.3	1.7	1.7±0.7	0.8	1.8	1.7	0.4	0.9	1.1±0.6
KLF6	Kruppel-like factor 6	1555832_s_at	19.6	20.0	26.0	18.3	16.7	20.1±3.5	34.8	33.1	47.3	36.8	37.0	37.8±5.5
	Kruppel-like factor 6	208960_s_at	2.7	1.8	2.7	1.2	2.1	2.1±0.6	3.2	3.0	4.9	5.1	5.0	4.2±1.1
	Kruppel-like factor 6	208961_s_at	4.6	2.1	5.0	4.3	5.4	4.3±1.3	6.0	6.4	7.0	6.3	10.6	7.2±1.9
	Kruppel-like factor 6	211610_at	2.3	0.9	1.5	1.5	1.4	1.5±0.5	1.9	2.6	1.9	1.3	1.5	1.8±0.5
	Kruppel-like factor 6	224606_at	26.6	19.0	25.6	24.4	35.0	26.1±5.7	48.4	46.2	53.4	46.8	45.3	48.1±3.2
KLF7 (UKLF)	Kruppel-like factor 7 (ubiquitous)	1555420_a_at	0.9	0.1	0.4	0.2	1.0	0.5±0.4	0.5	1.0	0.8	0.7	0.6	0.7±0.2
	Kruppel-like factor 7 (ubiquitous)	204334_at	0.9	0.7	0.7	1.4	1.3	1.0±0.3	1.5	1.2	2.4	1.6	1.9	1.7±0.5
	Kruppel-like factor 7 (ubiquitous)	238482_at	0.6	0.4	1.0	1.2	1.3	0.9±0.4	0.8	1.1	1.1	1.2	1.6	1.2±0.3
KLF8	Kruppel-like factor 8	219930_at	0.6	1.6	0.1	0.1	0.1	0.5±0.6	0.9	1.8	0.3	0.5	0.6	0.8±0.6
	Kruppel-like factor 8	230986_at	0.1	0.3	0.6	0.1	0.4	0.3±0.2	0.0	0.3	0.2	0.5	0.1	0.2±0.2
KLF9	Kruppel-like factor 9	203541_s_at	1.4	1.5	1.0	0.8	1.3	1.2±0.3	0.7	1.4	1.0	0.7	0.8	0.9±0.3
	Kruppel-like factor 9	203542_s_at	54.8	42.5	55.0	44.4	56.6	50.7±6.6	59.8	56.9	54.5	52.3	49.6	54.6±4.0
	Kruppel-like factor 9	203543_s_at	35.5	25.3	33.6	28.4	34.4	31.4±4.4	34.2	30.8	34.9	28.5	31.9	32.1±2.6
	Kruppel-like factor 9	228474_s_at	1.2	0.8	1.6	0.3	0.2	0.8±0.6	0.2	0.1	0.5	0.5	0.3	0.3±0.2
	Kruppel-like factor 9	230636_s_at	1.1	0.8	1.8	1.0	1.0	1.2±0.4	0.7	0.5	0.2	1.6	1.0	0.8±0.5
KLF10	Kruppel-like factor 10	202393_s_at	55.2	44.8	60.2	54.3	63.2	55.5±7.0	69.7	59.5	67.1	66.0	67.3	65.9±3.8

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KLF11	Kruppel-like factor 11	1553137_s_at	0.2	1.6	0.3	2.2	1.9	1.2±0.9	1.6	1.0	2.3	0.3	0.3	1.1±0.9
	Kruppel-like factor 11	218486_at	3.9	3.6	4.8	5.9	4.8	4.6±0.9	3.7	4.9	3.1	4.8	2.8	3.8±1.0
KLF12	Kruppel-like factor 12	206965_at	0.4	0.6	0.5	0.5	0.1	0.4±0.2	0.8	0.4	0.8	0.5	0.5	0.6±0.2
	Kruppel-like factor 12	206966_s_at	1.9	1.6	2.9	0.3	1.8	1.7±0.9	0.3	1.0	1.7	0.5	0.6	0.8±0.6
	Kruppel-like factor 12	208467_at	0.2	1.0	0.2	0.1	0.1	0.3±0.4	0.1	1.0	0.6	0.6	0.4	0.5±0.3
	Kruppel-like factor 12	214276_at	0.2	0.2	0.2	0.9	0.7	0.5±0.3	0.1	0.1	0.6	1.6	0.2	0.5±0.6
	Kruppel-like factor 12	227261_at	0.2	1.8	1.3	2.3	1.2	1.4±0.8	0.7	1.2	1.8	0.9	2.5	1.4±0.7
	Kruppel-like factor 12	229881_at	0.1	0.1	0.1	0.3	0.2	0.2±0.1	0.1	0.8	0.2	0.3	0.1	0.3±0.3
	Kruppel-like factor 12	238940_at	0.2	0.0	0.1	0.6	1.0	0.4±0.4	0.1	0.1	0.2	0.8	0.1	0.3±0.3
	Kruppel-like factor 12	239019_at	0.1	0.1	0.1	0.1	0.1	0.1±0.0	0.1	0.2	0.1	0.4	0.1	0.2±0.1
KLF13	Kruppel-like factor 13	219878_s_at	5.6	3.5	3.0	2.4	8.6	4.6±2.5	2.0	2.7	1.4	3.8	3.2	2.6±1.0
	Kruppel-like factor 13	225390_s_at	25.1	25.5	40.2	26.3	32.1	29.8±6.4	20.8	18.1	24.7	20.3	17.7	20.3±2.8
KLF14	Kruppel-like factor 14	1552813_at	0.3	0.2	0.7	0.2	0.3	0.4±0.2	0.3	1.7	1.2	0.5	1.2	1.0±0.6
	Kruppel-like factor 14	1552814_a_at	0.3	0.1	0.7	0.1	0.5	0.3±0.3	0.2	1.2	0.1	0.6	1.0	0.6±0.5
KLF15	Kruppel-like factor 15	221302_at	2.5	1.2	0.9	0.1	0.2	0.1±0.1	0.2	0.5	0.3	0.2	0.4	0.3±0.1
	Kruppel-like factor 15	231015_at	3.7	2.7	3.8	3.1	3.5	3.4±0.5	3.3	2.0	2.9	3.1	2.5	2.8±0.5
KLF16	Kruppel-like factor 16	224111_x_at	0.5	0.1	0.2	0.3	0.1	0.3±0.2	0.1	0.2	0.5	0.3	0.2	0.3±0.1
	Kruppel-like factor 16	226328_at	0.3	0.1	0.1	0.2	0.4	0.2±0.1	0.5	0.1	0.1	1.1	0.3	0.4±0.4
KLF17	Kruppel-like factor 17	1553891_at	0.2	0.2	1.0	0.6	0.2	0.4±0.4	0.4	0.3	0.2	0.2	0.2	0.3±0.1
NFE2 (p45 NF-E2)	nuclear factor (erythroid-derived 2), 45kDa	209930_s_at	74.3	63.2	72.5	68.1	75.6	70.7±5.1	60.2	61.6	66.7	54.1	43.6	57.2±8.8
MAFK (p18 NF-E2)	v-maf musculoaponeurotic fibrosarcoma oncogene homolog K (avian)	206750_at	1.1	1.0	1.1	1.0	1.3	1.1±0.1	1.3	0.2	0.8	0.6	1.2	0.8±0.4
	v-maf musculoaponeurotic fibrosarcoma oncogene homolog K (avian)	226206_at	2.4	2.1	2.5	2.1	3.2	2.5±0.5	3.3	2.9	3.5	1.7	2.9	2.9±0.7
NR2F2 (NF-E3, COUP-TFII)	nuclear receptor subfamily 2, group F, member 2	209119_x_at	0.1	0.1	0.1	0.2	0.0	0.1±0.1	0.8	0.1	0.1	0.1	0.1	0.2±0.3
	nuclear receptor subfamily 2, group F, member 2	209120_at	0.5	0.0	0.1	0.6	0.2	0.3±0.2	0.7	0.1	0.3	1.0	0.1	0.4±0.4
	nuclear receptor subfamily 2, group F, member 2	209121_x_at	0.4	0.8	0.2	0.8	0.9	0.6±0.3	0.2	1.6	0.2	0.3	0.1	0.7±0.6
	nuclear receptor subfamily 2, group F, member 2	215073_s_at	0.0	0.3	0.2	0.2	0.1	0.1±0.1	0.3	0.8	0.2	0.0	0.3	0.3±0.3
	nuclear receptor subfamily 2, group F, member 2	229092_at	0.1	0.1	0.5	0.7	0.8	0.4±0.3	0.8	0.3	0.1	0.1	0.6	0.3±0.3
NR2F1 (COUP-TFI)	nuclear receptor subfamily 2, group F, member 1	209505_at	0.1	0.5	0.1	0.1	0.1	0.2±0.2	0.5	0.7	0.4	0.8	0.0	0.5±0.3
	nuclear receptor subfamily 2, group F, member 1	209506_s_at	1.6	0.3	0.7	1.6	1.1	1.1±0.6	1.7	1.5	1.4	1.5	0.3	1.3±0.6
NF-E4	transcription factor NF-E4	1560527_at	0.0	0.1	0.2	0.1	0.1	1.1±0.0	0.7	0.2	0.2	0.3	0.2	0.3±0.2
BCL11A	B-cell CLL/lymphoma 11A (zinc finger protein)	1559078_at	14.6	4.9	8.6	7.3	9.4	9.0±3.6	7.3	5.7	5.5	4.8	6.1	5.9±0.9
	B-cell CLL/lymphoma 11A (zinc finger protein)	210347_s_at	6.3	6.0	6.8	6.5	10.9	7.3±2.0	6.2	5.8	4.7	4.6	4.8	5.2±0.7
	B-cell CLL/lymphoma 11A (zinc finger protein)	219497_s_at	45.8	32.9	38.9	34.5	34.7	37.3±5.2	21.8	21.1	19.2	17.5	18.6	19.6±1.8
	B-cell CLL/lymphoma 11A (zinc finger protein)	219498_s_at	29.3	23.4	24.3	21.4	24.7	24.6±2.9	14.2	13.0	10.5	9.7	9.6	11.4±2.1
	B-cell CLL/lymphoma 11A (zinc finger protein)	222891_s_at	59.1	47.1	53.5	45.8	54.6	52.0±5.5	31.3	30.4	22.5	29.5	27.4	28.2±3.5
HBS1L	HBS1-like (S. cerevisiae)	209314_s_at	41.3	36.0	32.2	31.4	33.4	34.8±4.0	39.9	25.6	25.0	25.8	25.8	28.4±6.4
	HBS1-like (S. cerevisiae)	209315_at	5.0	2.9	4.0	4.5	4.9	4.3±0.9	4.7	3.9	2.8	3.6	6.3	4.3±1.3
	HBS1-like (S. cerevisiae)	209316_s_at	87.6	55.3	69.5	58.7	61.8	66.6±12.9	62.7	56.6	59.8	46.7	48.8	54.9±6.9
	HBS1-like (S. cerevisiae)	240602_at	1.4	3.8	2.2	2.6	2.4	2.5±0.9	1.8	1.2	2.0	2.1	2.6	2.0±0.5
IKZF1	IKAROS family zinc finger 1 (Ikaros)	1565816_at	0.2	0.1	0.2	0.0	0.1	0.1±0.1	0.1	0.0	0.1	0.0	0.0	0.1±0.0
	IKAROS family zinc finger 1 (Ikaros)	1565817_at	0.5	0.4	0.1	1.6	0.8	0.7±0.5	0.2	1.5	0.7	2.1	0.2	0.9±0.9
	IKAROS family zinc finger 1 (Ikaros)	1565818_s_at	1.1	1.4	1.7	1.0	0.6	1.2±0.4	1.4	0.1	0.4	1.4	1.4	0.9±0.6
	IKAROS family zinc finger 1 (Ikaros)	205038_at	5.2	4.3	4.1	5.0	3.8	4.5±0.6	3.0	3.3	6.8	4.6	3.6	4.3±1.5
	IKAROS family zinc finger 1 (Ikaros)	205039_s_at	13.4	14.1	19.0	15.8	19.8	16.4±2.9	11.3	11.7	11.5	8.8	10.9	10.8±1.1
	IKAROS family zinc finger 1 (Ikaros)	216901_s_at	1.1	0.2	0.9	1.4	0.7	0.9±0.4	0.5	0.3	1.5	0.5	0.1	0.6±0.6

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			#1	#2	#3	#4	#5	Ave±SD	#1	#2	#3	#4	#5	Ave±SD
	IKAROS family zinc finger 1 (Ikaros)	220704_at	2.6	2.4	1.2	2.1	2.5	2.2±0.6	2.1	1.9	2.2	2.1	1.3	1.9±0.4
	IKAROS family zinc finger 1 (Ikaros)	227344_at	8.0	5.1	6.1	4.6	7.4	6.2±1.5	4.0	3.9	5.2	5.2	3.1	4.3±0.9
	IKAROS family zinc finger 1 (Ikaros)	227346_at	24.4	22.8	32.3	24.6	26.2	26.1±3.7	17.4	14.4	17.5	16.5	17.3	16.6±1.3
MYB	v-myb myeloblastosis viral oncogene homolog (avian)	204798_at	52.3	39.5	46.0	60.4	55.1	50.7±8.1	43.7	33.5	33.2	51.8	45.0	41.5±8.0
	v-myb myeloblastosis viral oncogene homolog (avian)	215152_at	1.5	0.8	0.1	1.8	1.6	1.2±0.7	1.2	1.0	1.3	1.7	1.5	1.3±0.3
NFYA	nuclear transcription factor Y, alpha	204107_at	6.9	2.4	5.8	5.1	4.4	4.9±1.7	4.4	3.0	5.3	3.7	3.5	4.0±0.9
	nuclear transcription factor Y, alpha	204108_at	6.7	6.2	6.3	7.4	8.6	7.0±1.0	8.6	8.6	8.3	9.9	8.1	8.7±0.7
	nuclear transcription factor Y, alpha	204109_s_at	3.3	2.9	4.4	4.1	4.6	3.9±0.7	4.6	6.7	5.6	4.7	3.9	5.1±1.1
	nuclear transcription factor Y, alpha	215720_s_at	1.7	0.3	0.9	3.1	2.5	1.7±1.1	1.4	2.2	1.8	2.1	2.1	1.9±0.3
NFYB	nuclear transcription factor Y, beta	218127_at	23.6	16.9	17.6	22.4	21.7	20.5±3.0	17.6	15.5	20.5	15.9	13.8	16.7±2.5
	nuclear transcription factor Y, beta	218128_at	11.1	7.7	10.7	12.7	13.6	11.2±2.3	7.4	7.4	7.9	7.1	7.7	7.5±0.3
	nuclear transcription factor Y, beta	218129_s_at	14.3	7.1	11.0	15.0	10.4	11.5±3.2	7.9	8.8	13.0	10.4	9.7	9.9±1.9
	nuclear transcription factor Y, beta	244704_at	2.5	1.7	2.2	2.8	1.6	2.1±0.5	1.4	1.7	1.8	1.7	2.5	1.8±0.4
POU2F1 (OCT-1)	POU class 2 homeobox 1	206789_s_at	3.9	2.9	4.5	3.8	4.3	3.9±0.6	4.0	4.0	3.7	3.4	3.7	3.7±0.3
	POU class 2 homeobox 1	227254_at	3.5	4.6	2.2	3.0	2.8	3.2±0.9	2.2	2.7	1.9	1.7	3.6	2.4±0.8
	POU class 2 homeobox 1	229753_at	1.5	1.6	0.4	0.9	1.0	1.1±0.5	0.8	0.1	1.0	1.3	1.0	0.8±0.4
	POU class 2 homeobox 1	244006_at	0.0	1.0	0.7	0.7	0.0	0.5±0.4	0.4	0.1	0.1	0.0	1.1	0.4±0.4
SOX6	SRY (sex determining region Y)-box 6	1563454_at	5.1	2.9	3.7	2.0	4.3	3.6±1.2	0.8	1.7	1.9	0.9	2.2	1.5±0.6
	SRY (sex determining region Y)-box 6	1570486_at	2.9	2.3	2.1	2.2	1.8	2.3±0.4	1.5	0.9	1.1	1.3	1.1	1.2±0.2
	SRY (sex determining region Y)-box 6	223865_at	2.1	1.8	1.5	2.3	3.0	2.2±0.6	0.2	1.2	0.2	0.2	0.7	0.5±0.4
	SRY (sex determining region Y)-box 6	224178_s_at	1.8	1.4	2.1	1.5	1.2	1.6±0.4	1.2	2.0	0.8	1.0	1.1	1.2±0.4
	SRY (sex determining region Y)-box 6	235526_at	12.8	8.1	11.2	11.6	13.1	11.4±2.0	9.5	8.3	8.3	10.1	13.0	9.9±1.9
SP1	Sp1 transcription factor	1553685_s_at	3.3	3.0	2.5	4.4	3.2	3.3±0.7	2.7	2.3	3.0	2.1	2.9	2.6±0.4
	Sp1 transcription factor	214732_at	2.0	0.7	1.4	0.2	0.6	1.0±0.7	1.1	0.5	2.4	1.0	1.7	1.3±0.7
	Sp1 transcription factor	224754_at	24.3	16.1	22.7	18.2	26.0	21.5±4.2	20.4	17.9	15.7	18.6	21.1	18.7±2.1
	Sp1 transcription factor	224760_at	15.1	11.0	13.3	13.3	15.9	13.7±1.9	12.8	9.8	13.5	13.7	9.7	11.9±2.0
TBP	TATA box binding protein	203135_at	10.4	8.5	9.0	9.2	10.9	9.6±1.0	6.9	9.0	9.6	7.4	9.2	8.4±1.2
YY1	YY1 transcription factor	200047_s_at	78.0	77.3	89.9	81.2	80.6	81.4±5.1	74.0	67.6	69.2	71.2	70.7	70.5±2.4
	YY1 transcription factor	201901_s_at	51.2	43.8	55.1	53.0	56.1	51.8±4.9	54.1	38.6	35.9	40.1	39.7	41.7±7.1
	YY1 transcription factor	201902_s_at	1.6	0.4	0.1	0.5	1.1	0.7±0.6	1.2	0.5	1.7	0.1	1.2	1.0±0.6
	YY1 transcription factor	213494_s_at	1.5	2.7	2.1	2.4	2.7	2.3±0.5	1.8	0.8	1.5	0.5	1.6	1.2±0.5
	YY1 transcription factor	224711_at	14.1	15.6	20.1	19.7	20.8	18.1±3.0	16.7	12.3	16.6	16.5	13.9	15.2±2.0
	YY1 transcription factor	224718_at	5.0	4.4	5.8	5.2	6.7	5.4±0.9	5.6	4.3	5.7	4.5	4.9	5.0±0.6
ZFPM1 (FOG-1)	zinc finger protein, multitype 1	237463_at	0.3	0.2	0.2	0.6	0.3	0.3±0.2	1.0	0.3	1.2	0.3	0.1	0.6±0.5
	zinc finger protein, multitype 1	242282_at	3.7	5.8	7.5	5.5	7.4	6.0±1.5	4.7	4.3	3.4	4.1	4.0	4.1±0.5