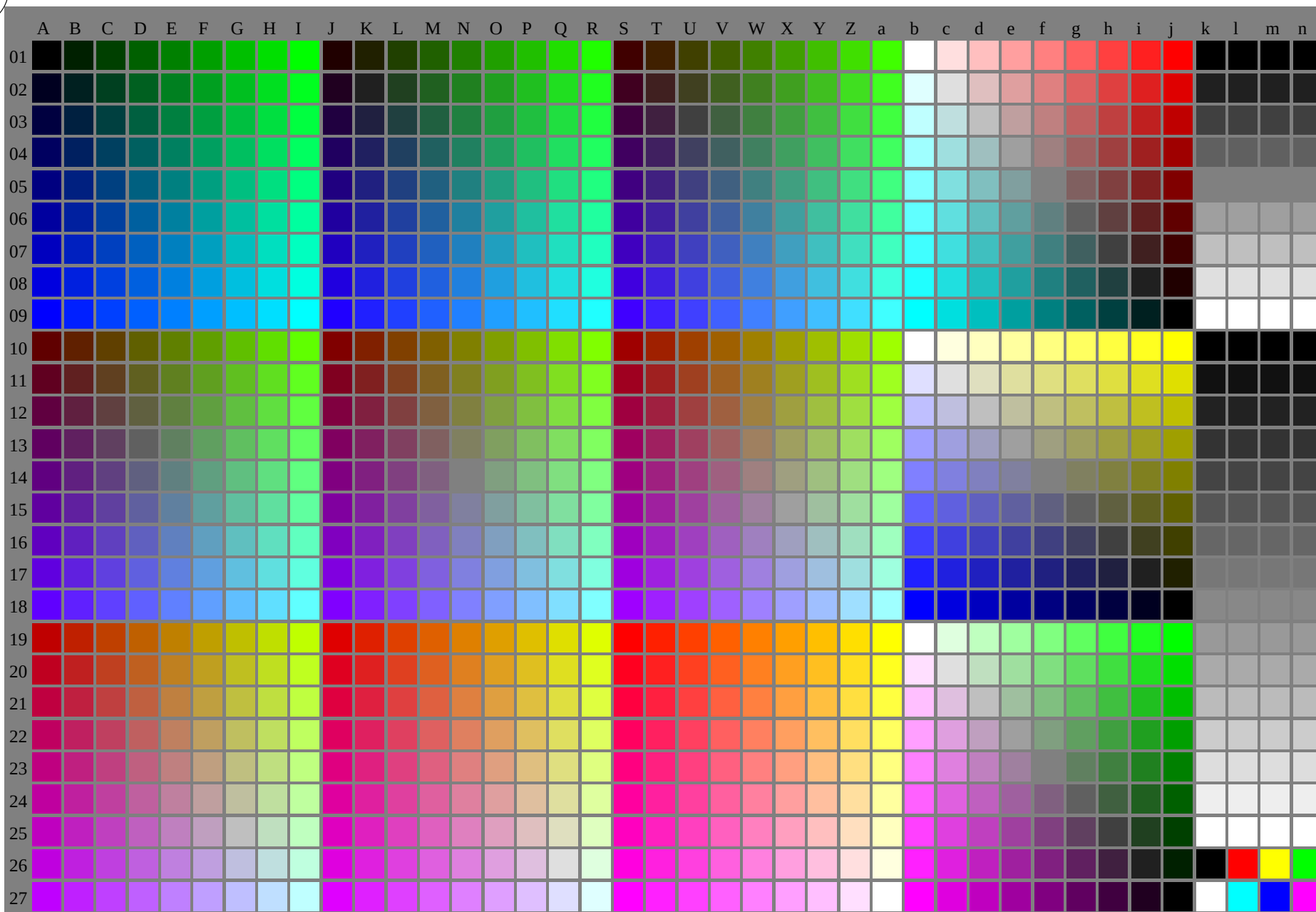


See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF> /.PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



LE520-7A, 1/16

TUB-test chart LE52; 1080 colours of system G; $Tr=0\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

input: *rgb setrgbcolor*
output: no change

See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF> / .PSS
Technical information: <http://www.ps-bam.de> or <http://130.149.60.45/~farbmehrlik>

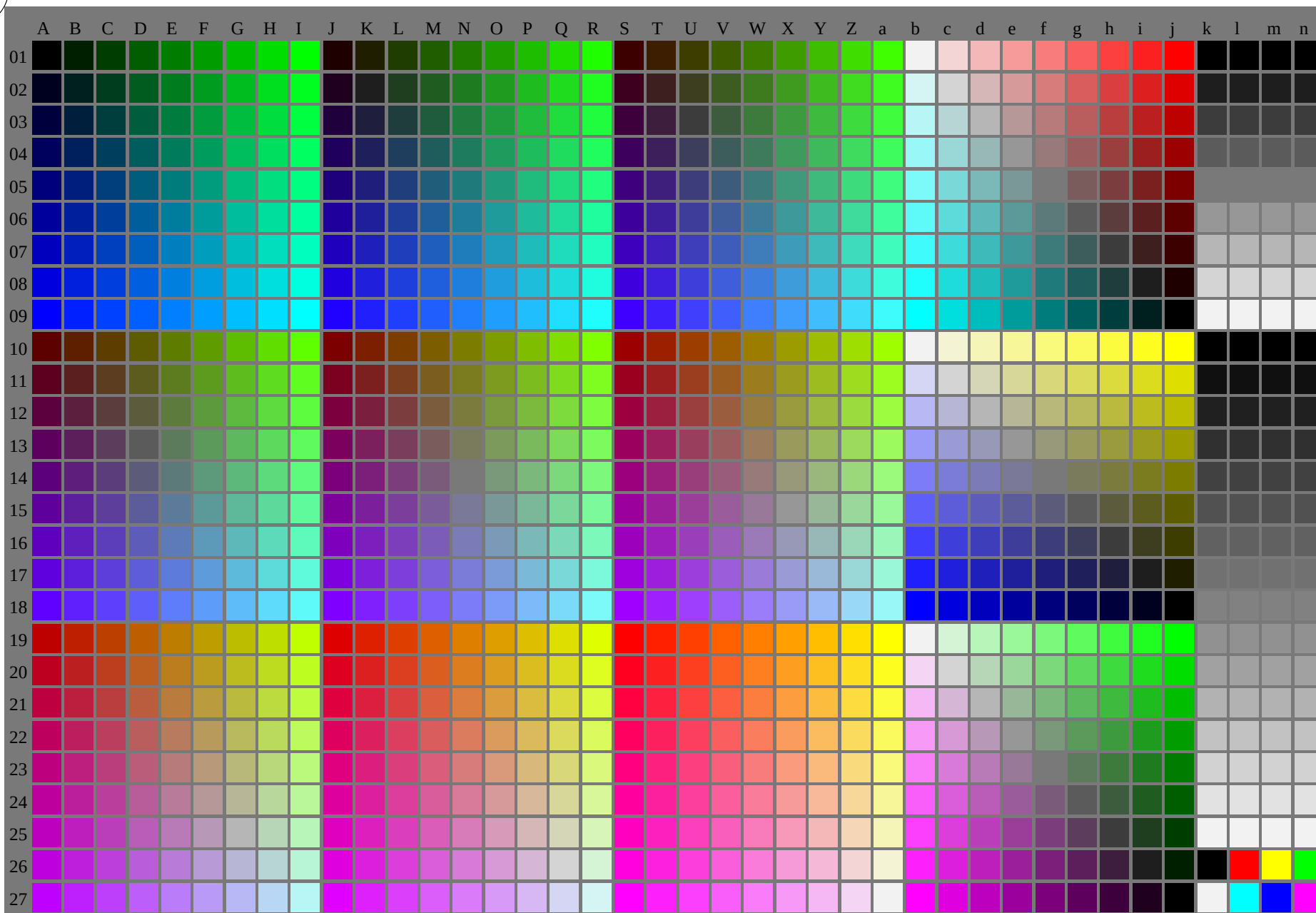
[illegible]

application for measurement of printer or monitor systems

TUB-test chart LE52; 1080 colours of system G; $Tr=0\%$, (Fadin) input:
Change of white to gray for sRGB display produces luminous colours output:

`rgb setrgbcolor`
t: no change

See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF /.PS>
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



LE520-7A, 3/16

TUB-test chart LE52; 1080 colours of system G; $Tr=2,5\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

input: *rgb setrgbcolor*
output: no change

Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

http://130.149.60.45/~farbmetrik/LE52/LE52L0NP.PDF /.PS; start output

N: No Output Linearization (OL) data in File (F), Startup (S) or Device (D), page 5/16

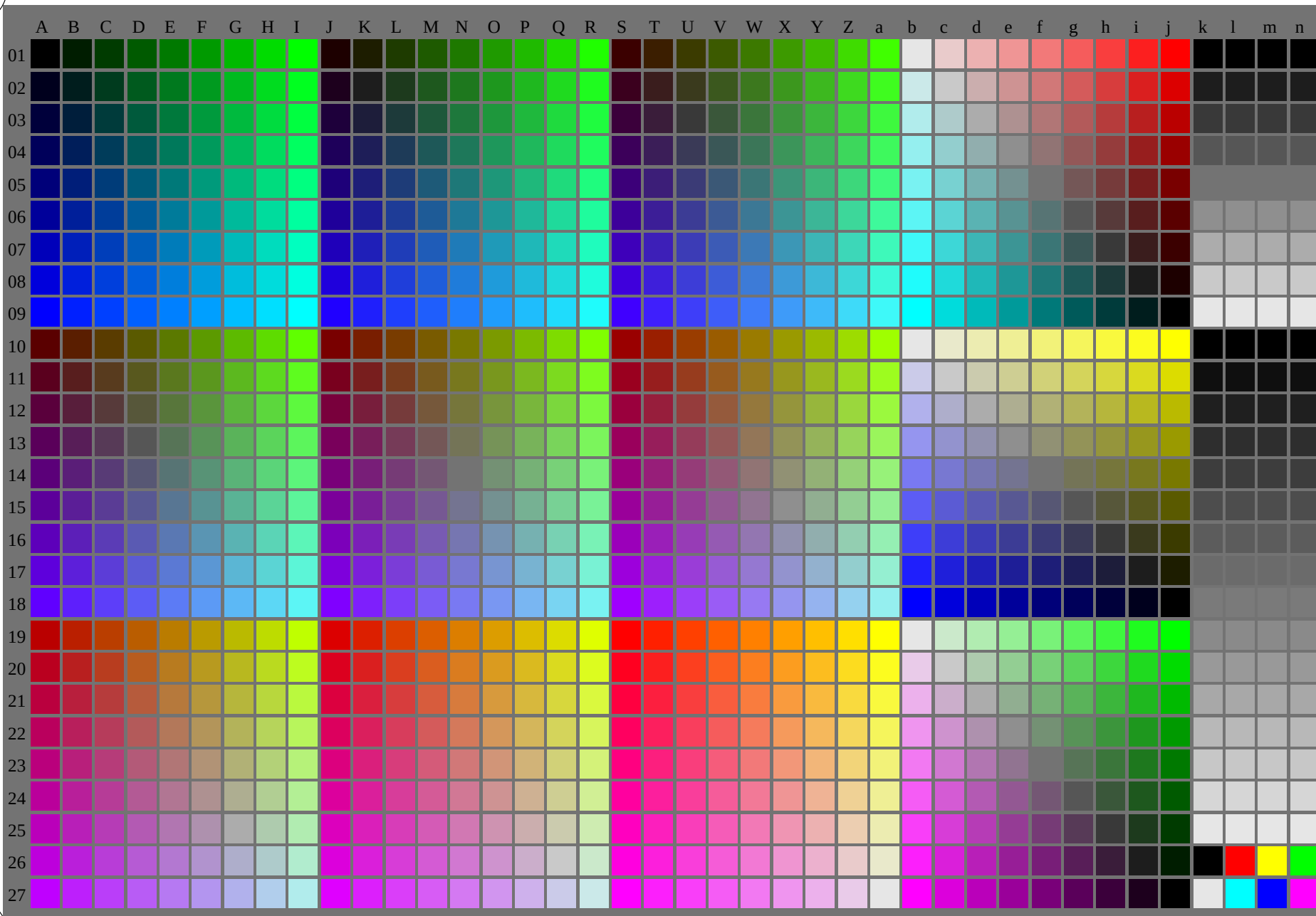
TUB-test chart LE52; 1080 colours of system G; $Tr=5\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

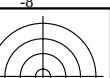
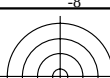
input: *rgb setrgbcolor*
output: no change

TUB registration: 20101101-LE52/LE52L0NP.PDF /.PS
application for measurement of printer or monitor systems

TUB material: code=rha4ta

See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF /.PS>
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>





See original or copy: <http://web.me.com/klaus-richter/LE52L/E52LONP.PDF> / .PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~fabneterik>

TUB registration: 20101101-LE52L/E52LONP.PDF / .PS
TUBmaterial: code=thafita

<http://130.149.60.45/~fabneterik/LE52L/E52LONP.PDF> / .PS; start output
N: No Output Linearization (CL) data in File (F), Startup (S) or Device (D), page 6/16

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	a	b	c	d	e	f	g	h	i	j	k	l	m	n																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																							
0000 A01	0009 B01	0018 C01	0027 D01	0036 E01	0045 F01	0054 G01	0063 H01	0072 I01	0081 J01	0090 K01	0100 L01	0109 M01	0118 N01	0126 O01	0135 P01	0144 Q01	0153 R01	0162 S01	0171 T01	0180 U01	0189 V01	0198 W01	0207 X01	0216 Y01	0225 Z01	0234 a01	0243 b01	0252 c01	0261 d01	0270 e01	0279 f01	0288 g01	0297 h01	0306 i01	0315 j01	0324 k01	0333 l01	0342 m01	0351 n01																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																								
0.0	0.114	0.231	0.351	0.475	0.601	0.731	0.864	1.0	0.0	0.0	0.0	0.0	0.0	0.114	0.231	0.351	0.475	0.601	0.731	0.864	1.0	0.0	0.0	0.115	0.231	0.351	0.475	0.601	0.731	0.864	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0

http://130.149.60.45/~farbmetrik/LE52/LE52L0NP.PDF /.PS; start output

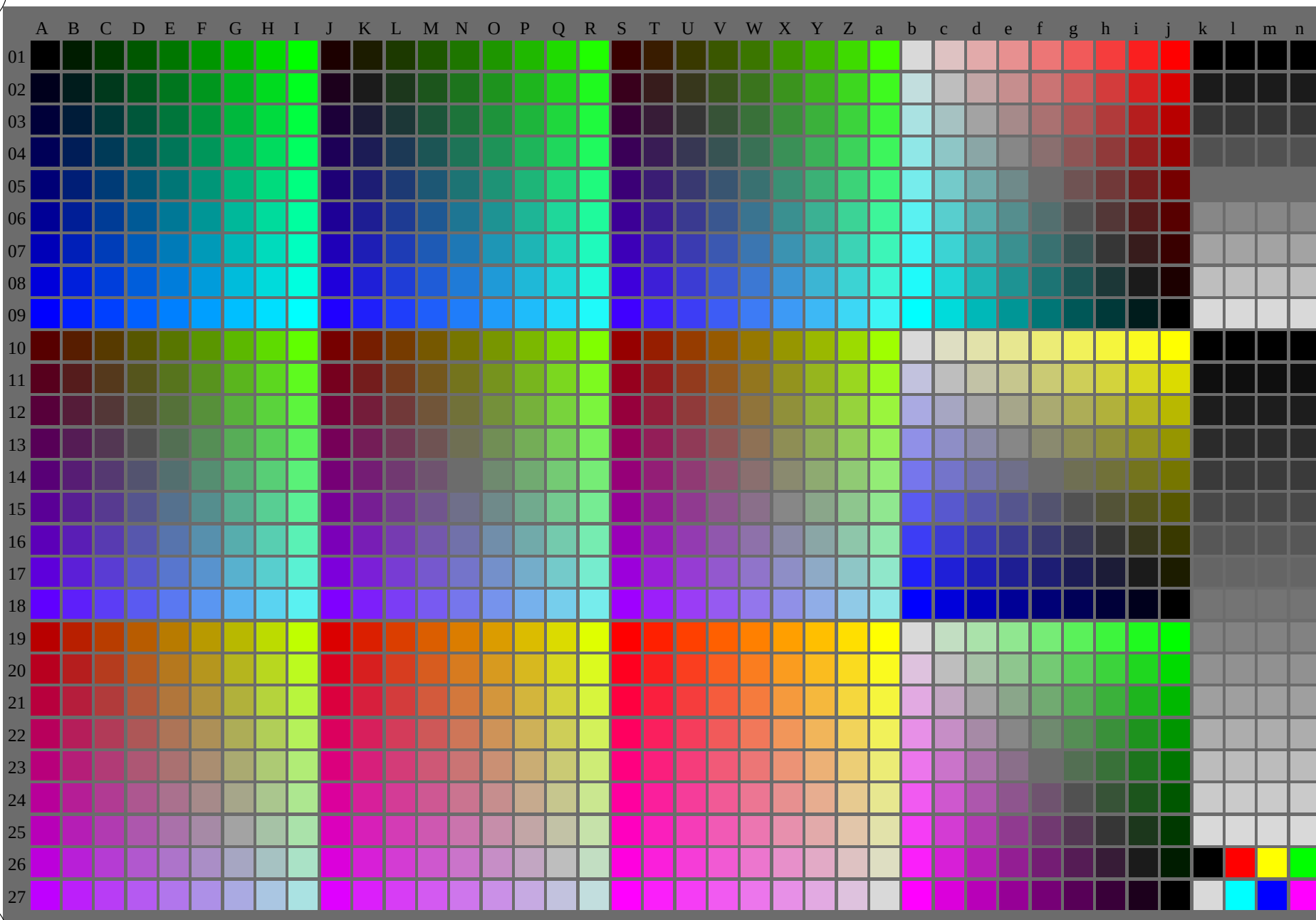
N: No Output Linearization (OL) data in File (F), Startup (S) or Device (D), page 7/16

TUB-test chart LE52; 1080 colours of system G; $Tr=7,5\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

input: *rgb setrgbcolor*
output: no change

TUB registration: 20101101-LE52/LE52L0NP.PDF /.PS
application for measurement of printer or monitor systems

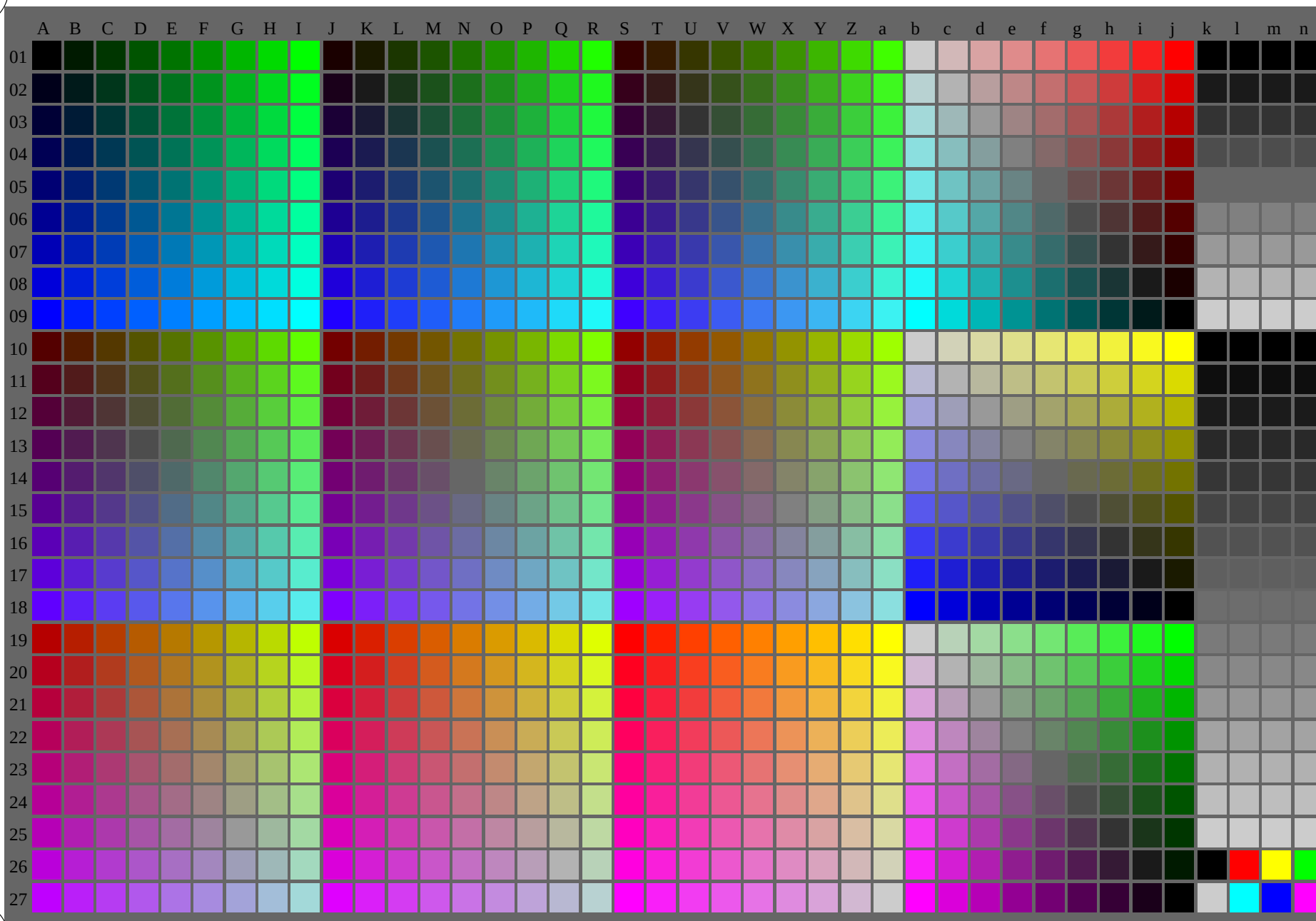
TUB material: code=rha4ta



See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF> /.PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

See original or copy: <http://web.me.com/klaus.richter/LE52LE52L0NP.PDF> / .PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF> /.PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



LE520-7A, 9/16

TUB-test chart LE52; 1080 colours of system G; $Tr=10\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

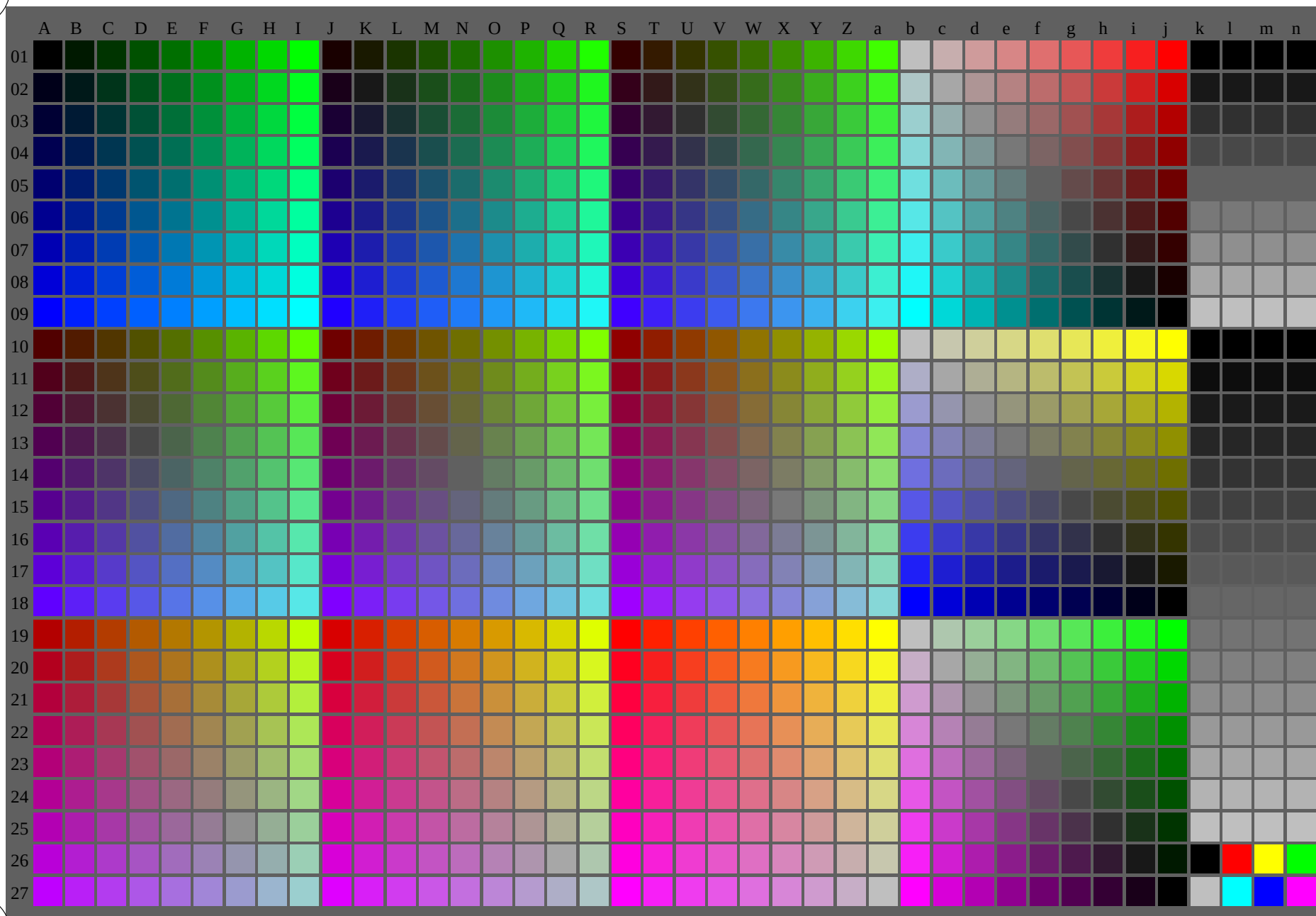
input: *rgb setrgbcolor*
output: no change

TUB registration: 20101101-LE52/LE52L0NP.PDF /.PS
application for measurement of printer or monitor systems

TUB material: code=rha4ta

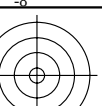
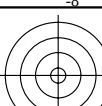
`rgb setrgbcolor`
 no change

See original or copy: <http://web.me.com/klaus.richter/LE52/LE52L0NP.PDF> /.PS
Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



TUB registration: 20101101-LE52/LE52L0NP.PDF /.PS
application for measurement of printer or monitor systems

TUB material: code=rha4ta



See original or copy: <http://web.me.com/klaus.richter/LE52L/LE52LONP.PDF> /PS
Technical information: <http://www.ps.pan.de> or <http://130.149.60.45/~fabnretik>

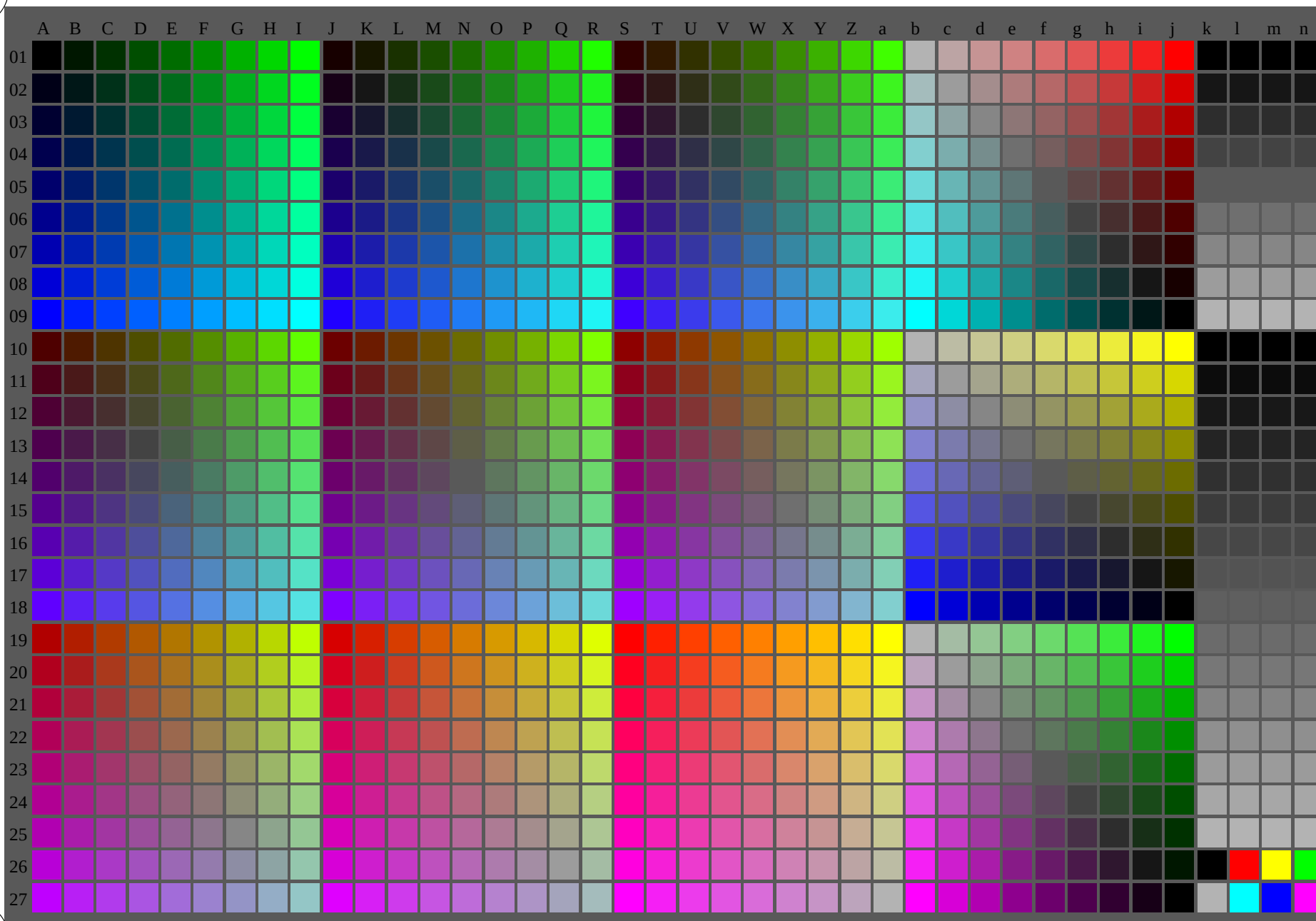
TUB registration: 2010101-LE52L/LE52LONP.PDF /PS
application for measurement of printer or monitor systems
TUBmaterial: code=ha4ta



<http://130.149.60.45/~fabnretik/LE52/LE52LONP.PDF> /PS; start output
N: No Output Linearization (OL) data in File (F), Startup (S) or Device (D), page 12/16

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	X	Y	Z	a	b	c	d	e	f	g	h	i	j	k	l	m	n
0000 A01	0009 B01	0018 C01	0027 D01	0036 E01	0045 F01	0054 G01	0063 H01	0072 I01	0081 J01	0090 K01	0099 L01	0108 M01	0117 N01	0126 O01	0135 P01	0144 Q01	0153 R01	0162 S01	0171 T01	0180 U01	0189 V01	0198 W01	0207 X01	0216 Y01	0225 Z01	0234 a01	0243 b01	0252 c01	0261 d01	0270 e01	0279 f01	0288 g01	0297 h01	0306 i01	0315 j01	0324 k01	0333 l01	0342 m01	0351 n01
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
0.000	0.007	0.010	0.013	0.016	0.019	0.022	0.025	0.028	0.031	0.034	0.037	0.040	0.043	0.046	0.049	0.052	0.055	0.058	0.061	0.064	0.067	0.070	0.073	0.076	0.079	0.082	0.085	0.088	0.091	0.094	0.097	0.100	0.103	0.106	0.109	0.112	0.115	0.118	0.121
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Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



LE520-7A, 13/16

TUB-test chart LE52; 1080 colours of system G; $Tr=15\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

input: *rgb setrgbcolor*
output: no change

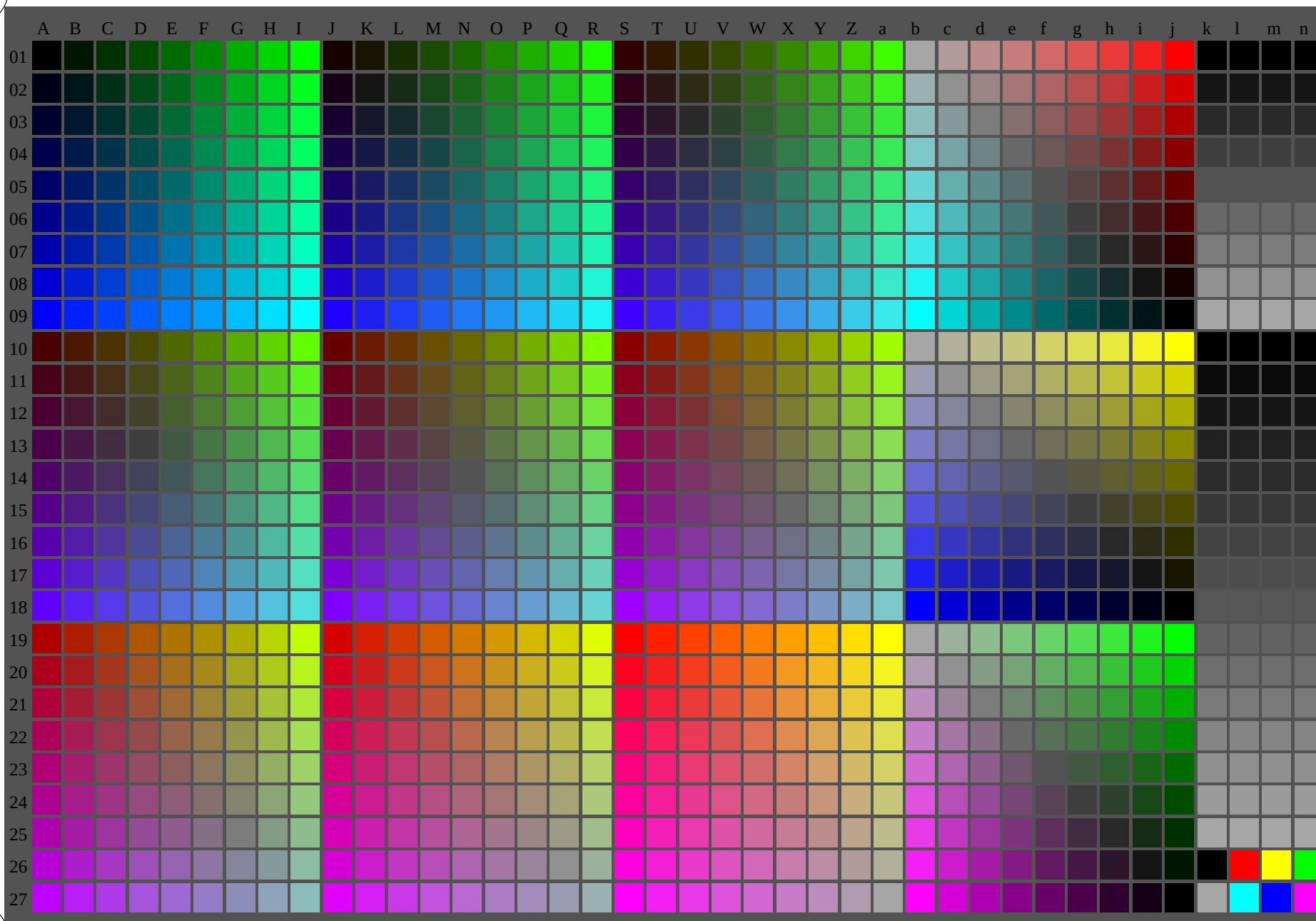
TUB registration: 20101101-LE52/LE52L0NP.PDF /.PS
application for measurement of printer or monitor systems

TUB material: code=rha4ta

TTTB material: code=rba4ta

[illegible]

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Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



LE520-7A, 15/16

TUB-test chart LE52; 1080 colours of system G; $Tr=17,5\%$, (Fadin)
Change of white to gray for sRGB display produces luminous colours

input: *rgb setrgbcolor*
output: no change

A diagram of a chromosome consisting of two sister chromatids. Each chromatid is represented by a vertical line. The top portion of each chromatid is colored pink and labeled 'M' (metacentromeric). The bottom portion is colored blue and labeled 'C' (centromeric). The two chromatids are joined at a central point labeled 'C', which represents the centromere. The ends of the chromatids are labeled 'M', representing the telomeres.

See original or copy: <http://web.me.com/klaus.richter/L.E52/L.E52L.0NP.PDF> , .PS
 Technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~fabmerrick>

TUB registration: 20101101-L-E52/L-E52L.0NP.PDF/.PS
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TUB material: code=rh4tra