

**Biomarker levels analysis according to crCR in the patients for whom pre-treatment formalin-fixed, paraffin-embedded tumor tissue block and cryopreserved tumor blocks were available (n=38)**

	<b>Response [95%CI]</b>		
	<b>Complete n=9</b>	<b>Incomplete n=29</b>	<b>All n=38</b>
<b>EGFR</b>			
Median (min, max)	0.6 [0.1;2]	0.7 [0.1;89.6]	0.6 [0.1;89.6]
Mean (SD)	0.8 (0.7)	4.1 (17.1)	3.3 (14.8)
n	9	27	36
<b>EGF</b>			
Median (min, max)	0.8 [0.1;3.3]	0.7 [0.1;76]	0.7 [0.1;76]
Mean (SD)	1.1 (1.1)	3.9 (14.5)	3.2 (12.5)
n	9	27	36
<b>TGF <math>\alpha</math></b>			
Median (min, max)	4.2 [1.6;15.4]	4.6 [0.6;33]	4.4 [0.6;33]
Mean (SD)	5.3 (4.1)	8.2 (8.9)	7.5 (8)
n	9	27	36
<b>HB-EGF</b>			
Median (min, max)	8.8 [0.6;28.9]	7.1 [0.7;66.9]	7.5 [0.6;66.9]
Mean (SD)	9.9 (9)	15.8 (17.7)	14.3 (16)
n	9	27	36
<b>BTC</b>			
Median (min, max)	3.6 [0.9;8.6]	3.1 [0;68.3]	3.2 [0;68.3]
Mean (SD)	3.9 (2.6)	6.6 (12.9)	5.9 (11.2)
n	9	27	36
<b>AREG</b>			
Median (min, max)	0.2 [0;3]	0.2 [0;92.9]	0.2 [0;92.9]
Mean (SD)	0.7 (1.1)	9.8 (24.8)	7.5 (21.8)
n	9	27	36

**EREG**

Median (min, max)	0 [0;0.9]	0 [0;9.5]	0 [0;9.5]
Mean (SD)	0.2 (0.3)	0.7 (1.9)	0.6 (1.7)
n	9	27	36

**Allele 1 intron 1 CA repeats**

Median (min, max)	16 [16;20]	16 [14;20]	16 [14;20]
Mean (SD)	16.7 (1.4)	16.4 (1.4)	16.5 (1.4)
n	9	27	36

**Allele 2 intron 1 CA repeats**

Median (min, max)	17 [16;20]	19 [15;22]	18 [15;22]
Mean (SD)	17.2 (1.4)	18.5 (2.2)	18.2 (2.1)
n	9	27	36

**EGFRvIII mutation**

No	5 (62.5%)	23 (88.5%)	28 (82.4%)
Yes	3 (37.5%)	3 (11.5%)	6 (17.7%)

**EGFR amplification**

No	3 (50%)	19 (82.6%)	22 (75.9%)
Yes	3 (50%)	4 (17.4%)	7 (24.1%)

EGF, epidermal growth factor; EGFR, epidermal growth factor receptor; TGF $\alpha$ , transforming growth factor  $\alpha$ ; HB-EGF, heparin-binding EGF-like growth factor; BTC, betacellulin; AREG, amphiregulin; EREG, epiregulin; SD, standard deviation