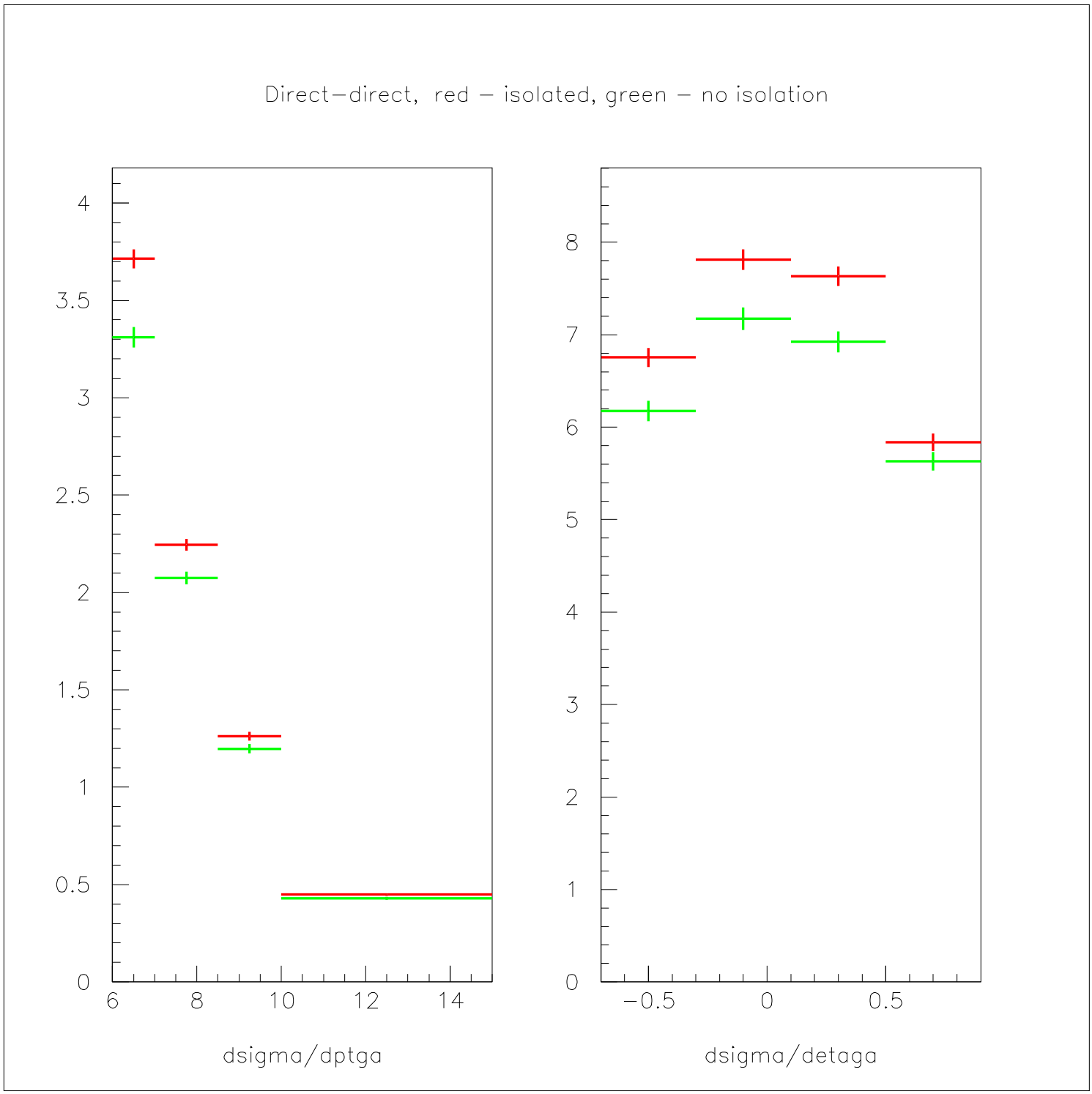
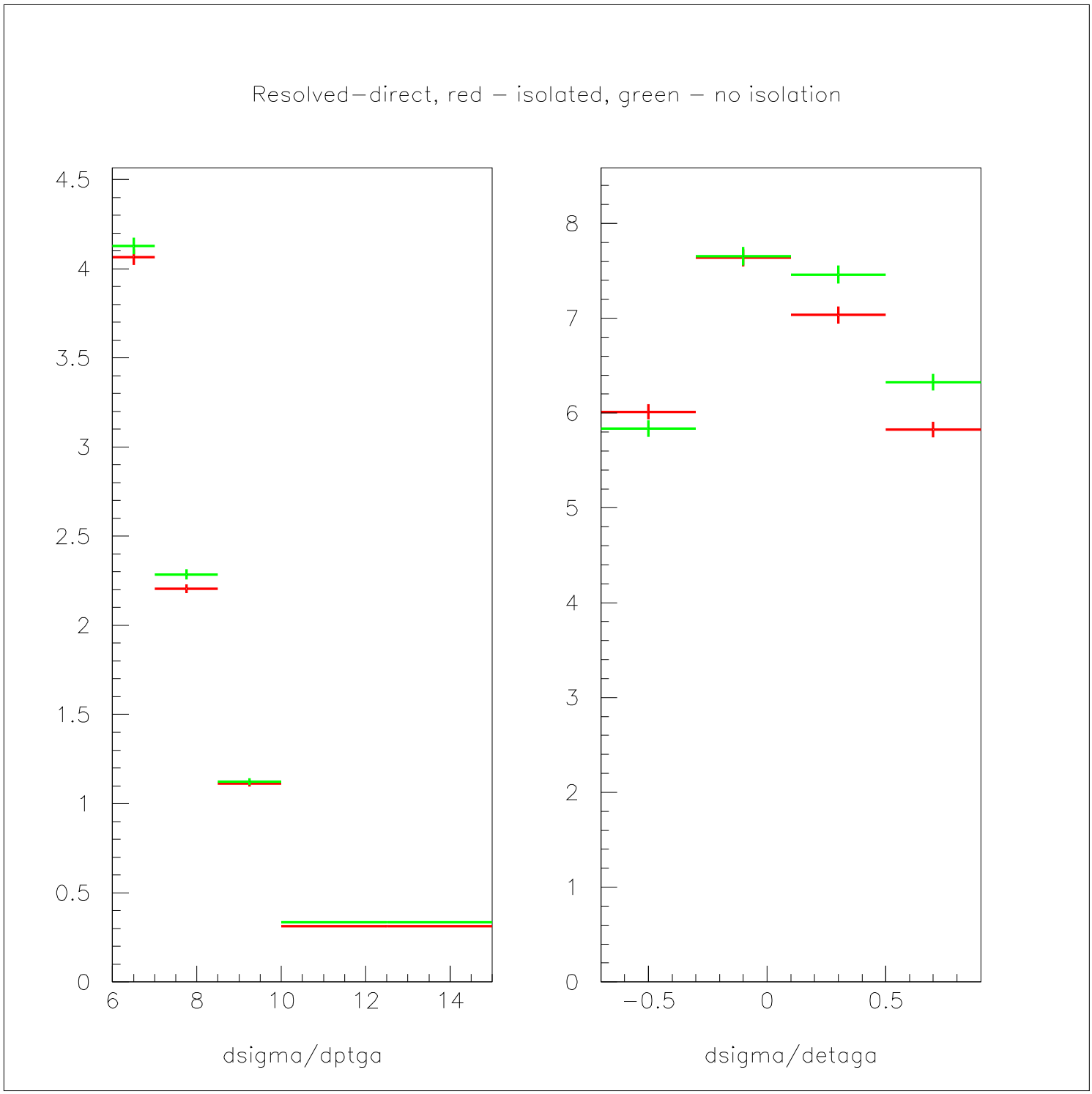
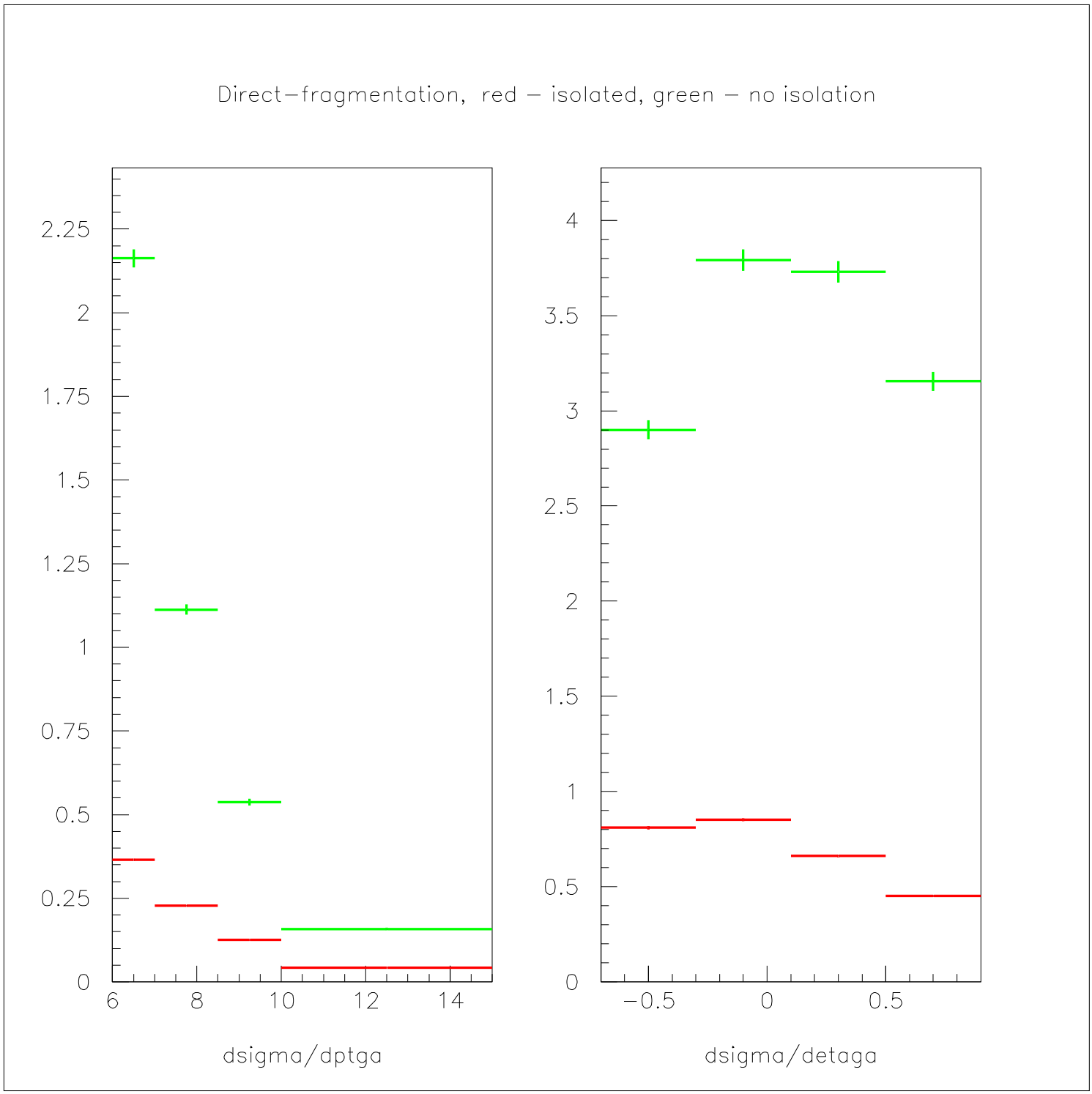
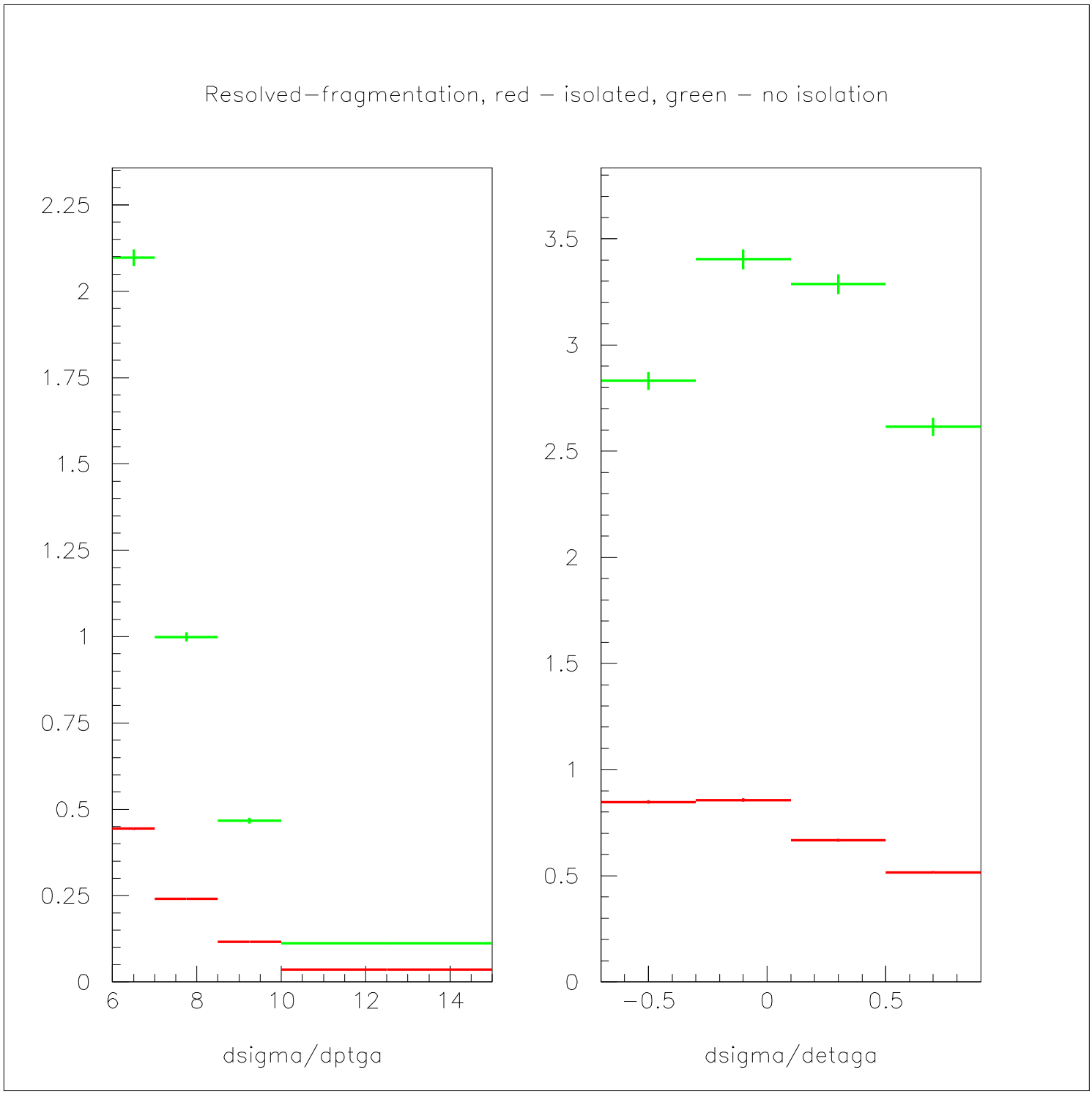

Direct−direct, red − isolated, green − no isolation
4
3.5
3
2.5
2
1.5
1
0.5
0
6
8
10
12
14
dsigma/dptga
8
7
6
5
4
3
2
1
0
−0.5
0
0.5
dsigma/detaga


Resolved−direct, red − isolated, green − no isolation
4.5
4
3.5
3
2.5
2
1.5
1
0.5
0
6
8
10
12
14
dsigma/dptga
8
7
6
5
4
3
2
1
0
−0.5
0
0.5
dsigma/detaga


Direct−fragmentation, red − isolated, green − no isolation
2.25
2
1.75
1.5
1.25
1
0.75
0.5
0.25
0
6
8
10
12
14
dsigma/dptga
4
3.5
3
2.5
2
1.5
1
0.5
0
−0.5
0
0.5
dsigma/detaga


Resolved−fragmentation, red − isolated, green − no isolation
2.25
2
1.75
1.5
1.25
1
0.75
0.5
0.25
0
6
8
10
12
14
dsigma/dptga
3.5
3
2.5
2
1.5
1
0.5
0
−0.5
0
0.5
dsigma/detaga