|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Antibody | Chase | % recognition  (mean) | | SD | | n | p |
| D7324 |  | Wt | M26P | Wt | M26P |  |  |
|  | 0’ | 105 | 84 | 65 | 45 | 3 | 0.669 |
|  | 30’ | 91 | 72 | 26 | 3 | 3 | 0.277 |
|  | 2h | 94 | 74 | 6 | 2 | 3 | 0.00541 |
| A32 |  |  |  |  |  |  |  |
|  | 0’ | 7 | 2 | 5 | 2 | 3 | 0.183 |
|  | 30’ | 13 | 4 | 5 | 3 | 3 | 0.0556 |
|  | 2h | 18 | 8 | 3 | 4 | 3 | 0.0257 |
| CD4-IgG |  |  |  |  |  |  |  |
|  | 0’ | 7 | 7 | 1 | 3 | 3 | 1.0 |
|  | 30’ | 43 | 27 | 9 | 5 | 3 | 0.0546 |
|  | 2h | 65 | 54 | 21 | 6 | 3 | 0.432 |

**Figure 7 – Source data 1:**

Autoradiographs from Figure 7F were quantified and each antibody normalized to the recognition of polyclonal antibody 40336 which recognizes all forms of gp160. Graphpad was used to calculate all statistical values for the above time points. p values were calculated using an unpaired, two-tailed t test with Welch’s correction.