

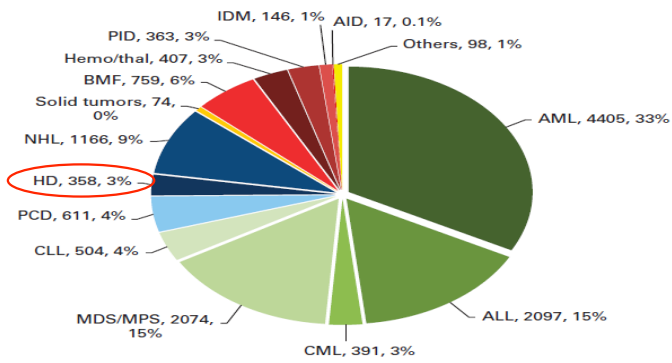
# Timing dell'allogtrapianto nel linfoma di Hodgkin nell'era dei nuovi farmaci

Bari, 6-7/6/2017

Luca Castagna

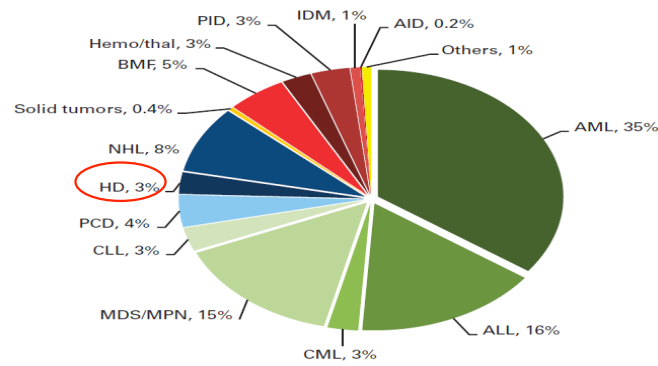
# Background

EBMT activity 2011

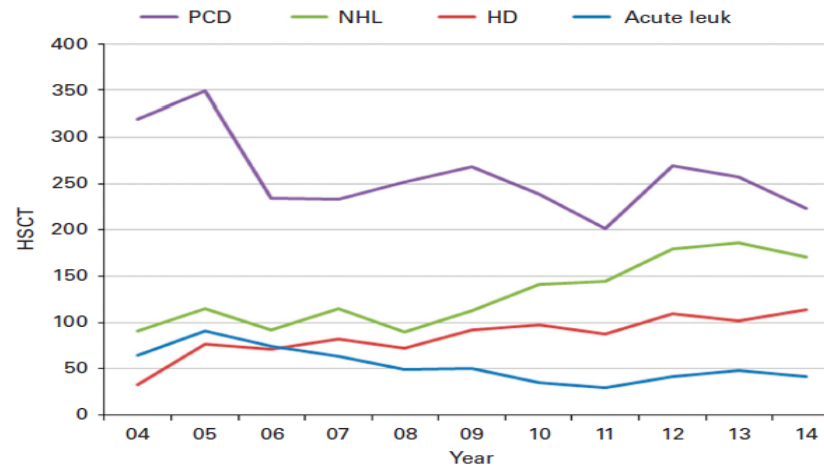
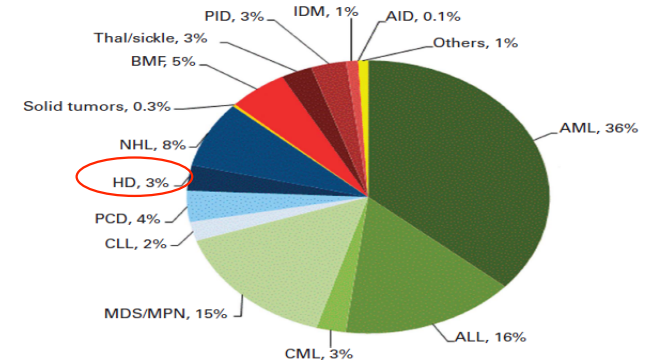


EBMT activity 2013

Allogeneic



EBMT activity 2014



# Background

| Disease status                       | HLAid sib | WM UD    | Alternative donor | ASCT      |
|--------------------------------------|-----------|----------|-------------------|-----------|
| <b>CR1</b>                           | GNR       | GNR      | GNR               | GNR       |
| <b>CT S relapse, no previous HDC</b> | D         | D        | GNR               | S         |
| <b>CT S relapse, previous HDC</b>    | <b>S</b>  | <b>S</b> | <b>CO</b>         | <b>CO</b> |
| <b>Refractory</b>                    | D         | D        | D                 | <b>CO</b> |

# HL and allo-SCT

- Does Graft versus HL exists?
  - **Probably yes => DLI effect**
- Are HL cells suitable of immunological control?
  - **Definitively yes => CPI effect**

# Competitors

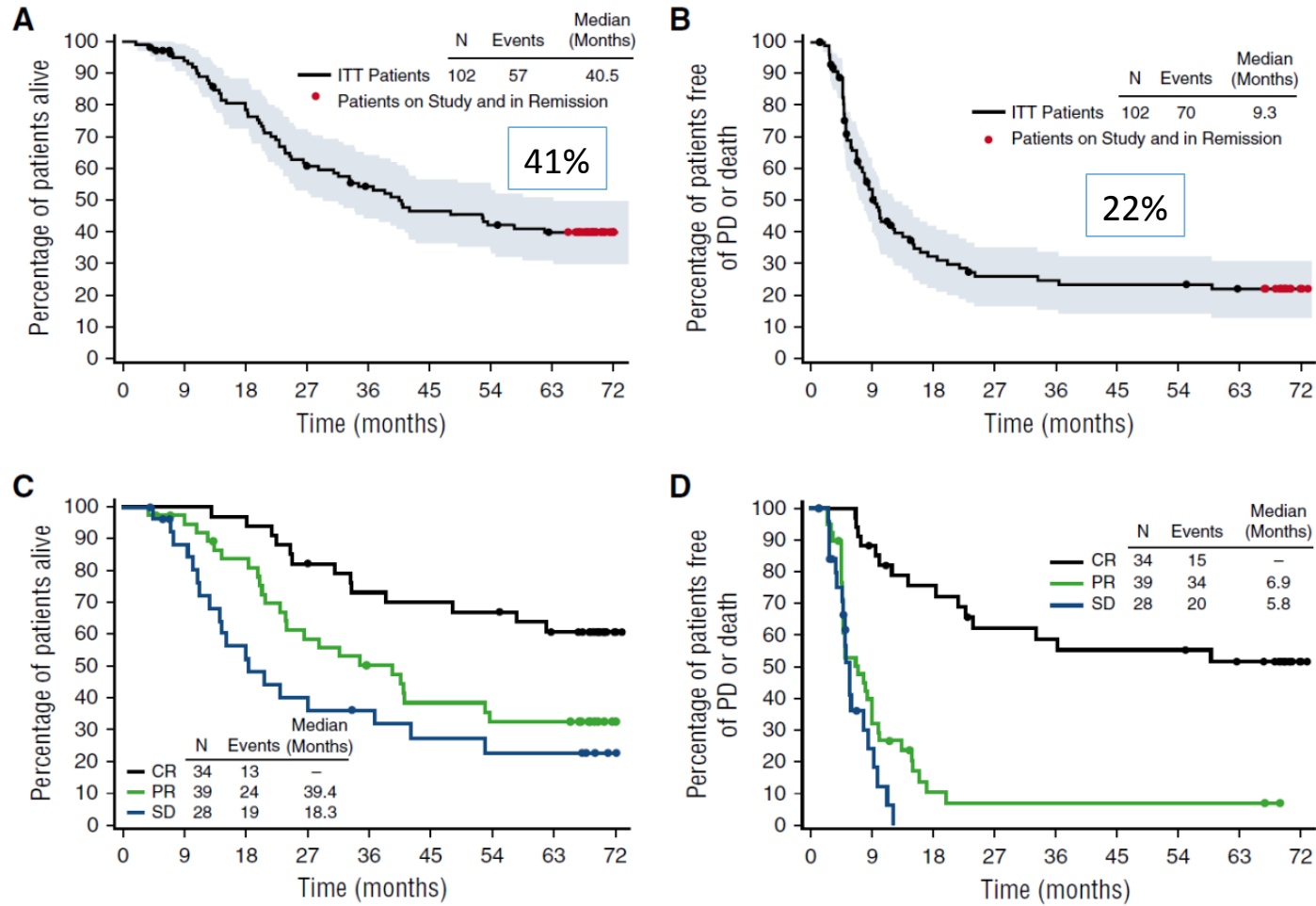
- Brentuximab vedotin (BV)
- Check point inhibitors (CPI)
  - Nivolumab
  - Pembrolizumab
- Association BV + CPI
- PI3Kinase inhibitors

## BV in relapsed/refractory patients

|                            | <b>Younes<br/>2012</b> | <b>Rothe<br/>2012</b> | <b>Zinzani<br/>2013</b> | <b>Gibb<br/>2013</b> |
|----------------------------|------------------------|-----------------------|-------------------------|----------------------|
| <b>N</b>                   | 102                    | 45                    | 65                      | 18                   |
| <b>Relapse after HDC</b>   | 100%                   | 87%                   | 92%                     | 33%                  |
| <b>ORR</b>                 | 75%                    | 60%                   | 29/70%                  | 72%                  |
| <b>CR</b>                  | 34%                    | 22%                   | 21%                     | 17%                  |
| <b>PR</b>                  | 41%                    | 38%                   | 8%                      | 55%                  |
| <b>DOR all responding</b>  | 6.7M                   | 8M                    | 6.8M                    | 5M                   |
| <b>DOR CR</b>              | 20 M                   | 13M (CR+PR)           | /                       | /                    |
| <b>ALLO (elegant/done)</b> | 102/6                  | 39/0                  | 62/9                    | 18/4                 |
| <b>OS</b>                  | 73%@3y                 | 83%@1y                | 74%@20M                 | /                    |
| <b>PFS</b>                 | 58%@3y                 | 43%@1y                | 23%@20M                 | 20%@1y               |
| <b>Response max</b>        | 3 cycles               | /                     | 3 cycles                | 4 cycles             |

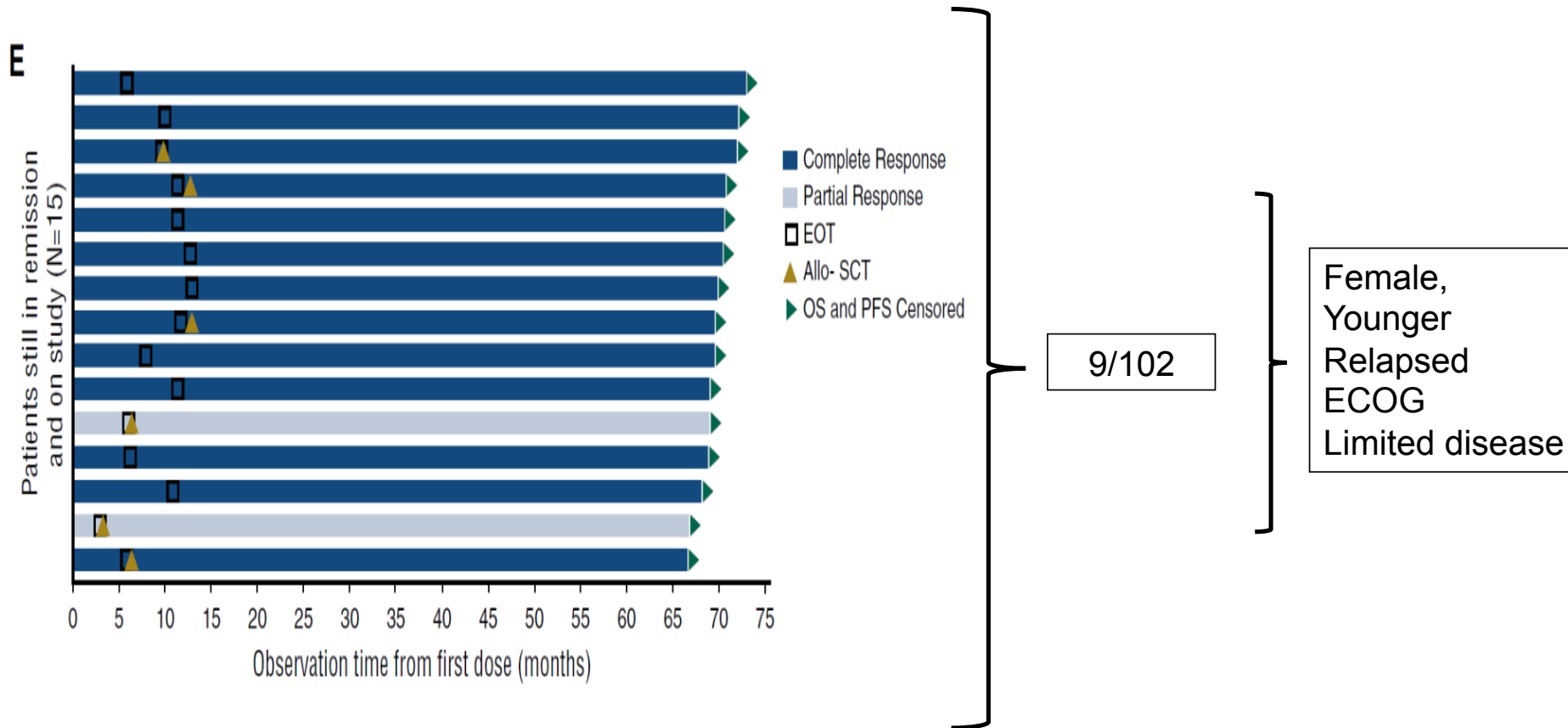
# Five-year survival and durability results of brentuximab vedotin in patients with relapsed or refractory Hodgkin lymphoma

Robert Chen,<sup>1,\*</sup> Ajay K. Gopal,<sup>2,\*</sup> Scott E. Smith,<sup>3</sup> Stephen M. Ansell,<sup>4</sup> Joseph D. Rosenblatt,<sup>5</sup> Kerry J. Savage,<sup>6</sup> Joseph M. Connors,<sup>6</sup> Andreas Engert,<sup>7</sup> Emily K. Larsen,<sup>8</sup> Dirk Huebner,<sup>9</sup> Abraham Fong,<sup>8</sup> and Anas Younes<sup>10</sup>



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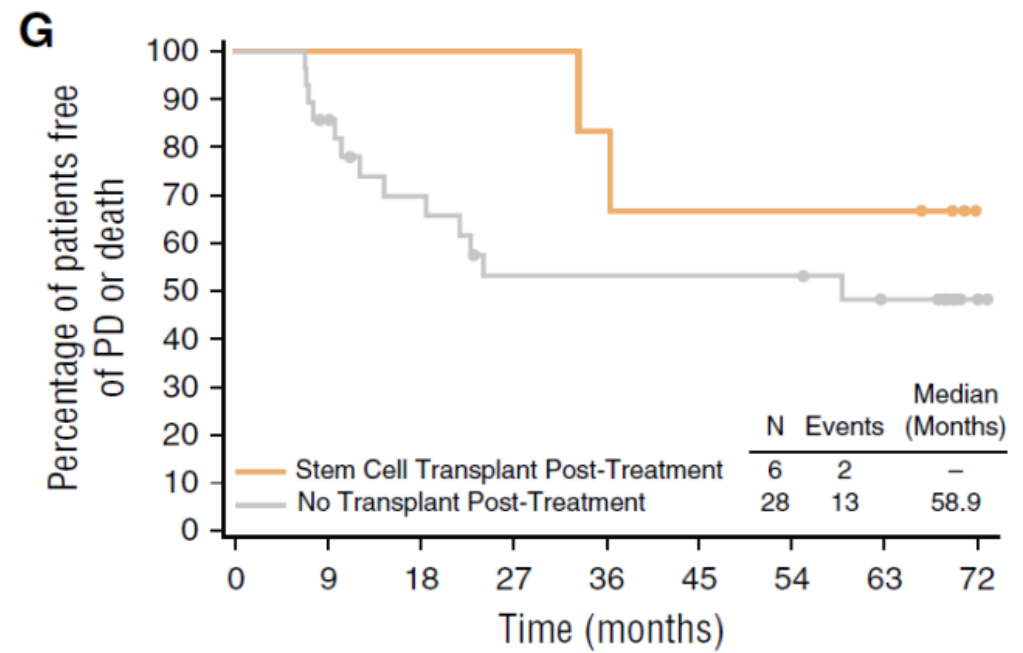
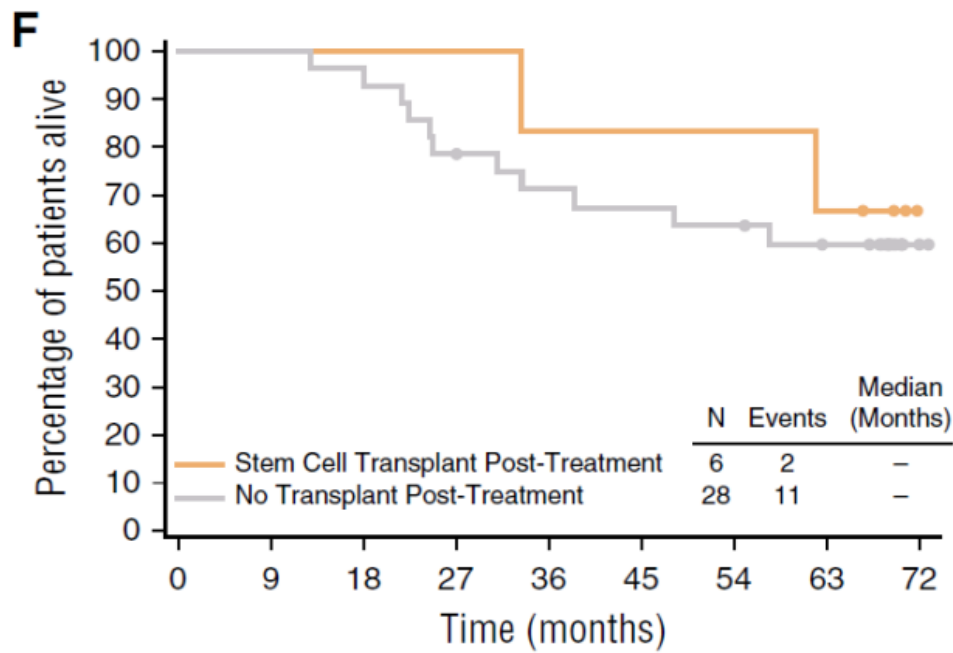
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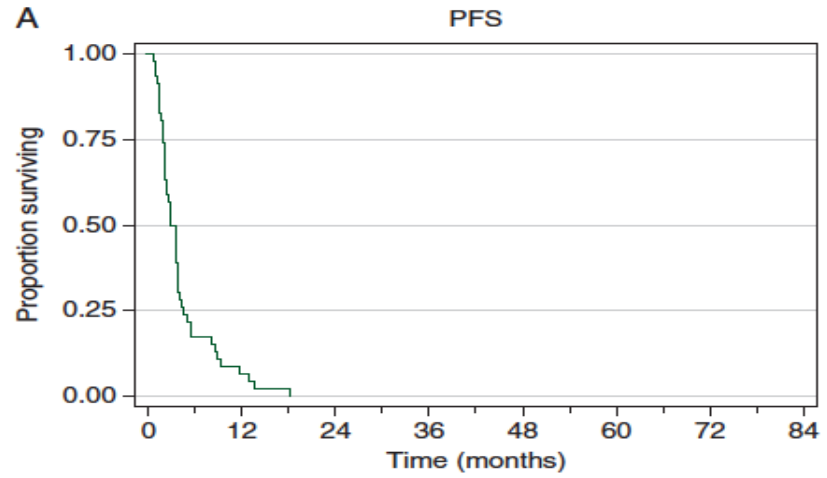


# Five-year survival and durability results of brentuximab vedotin in patients with relapsed or refractory Hodgkin lymphoma

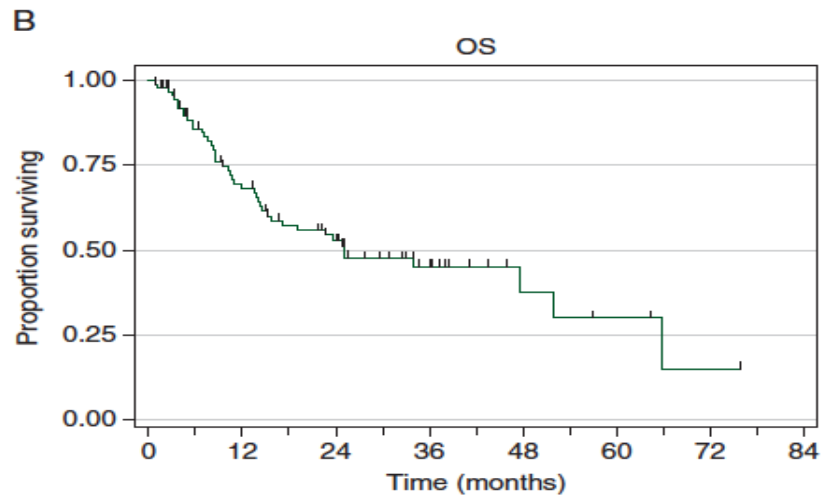
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# BV in relapsed/refractory patients



PFS 3.5 M

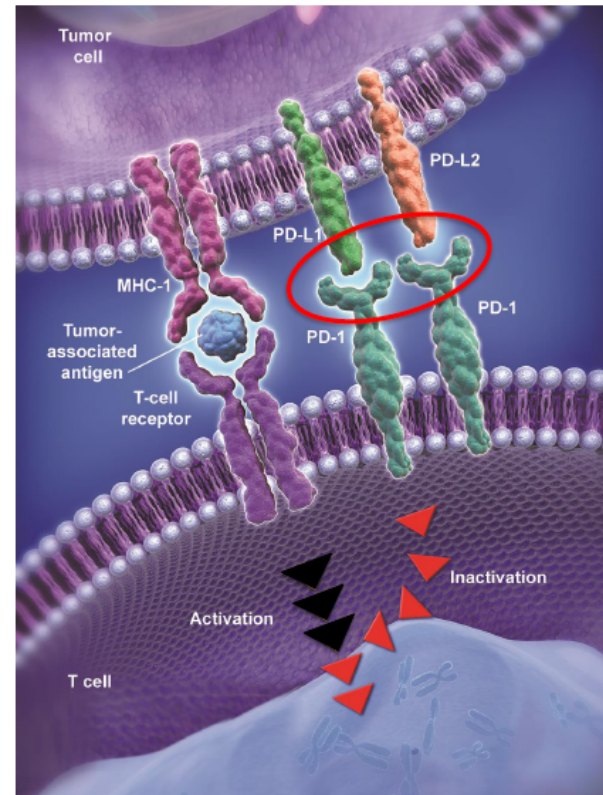


OS 24M

n = 100, 71% HDC

# The PD-1 and PD-L1/L2 Pathway

- PD-1 is an immune checkpoint receptor
- **Binding of PD-1 by its ligands PD-L1 or PD-L2 leads to downregulation of T-cell function**
- **This mechanism is usurped by many tumors**

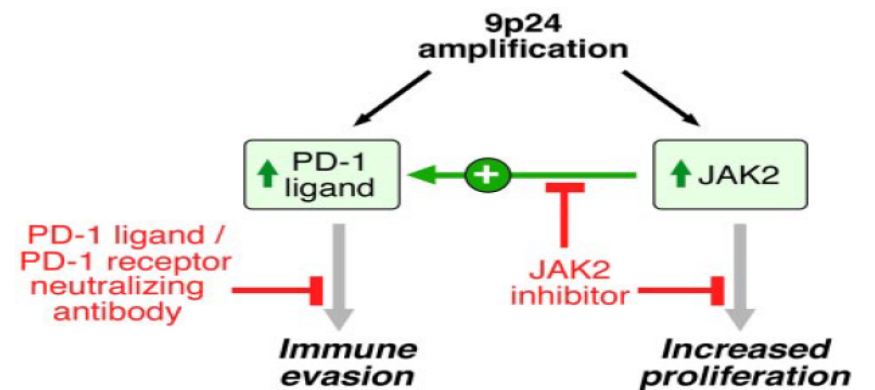
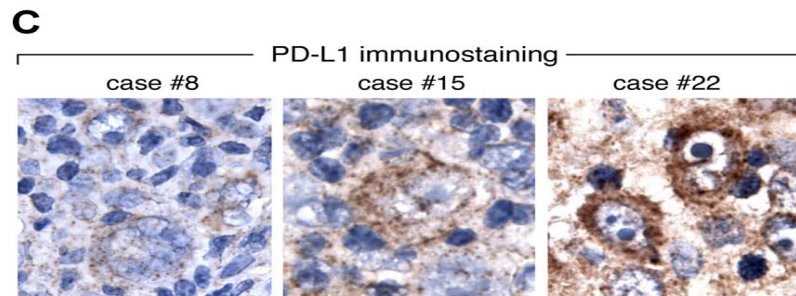
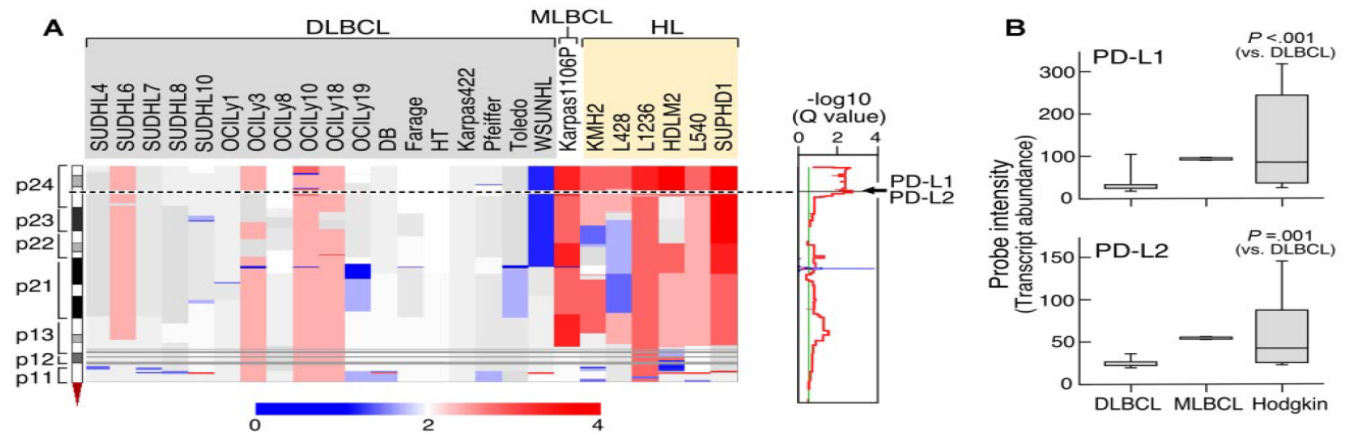


Keir ME et al. *Annu Rev Immunol.* 2008.  
Pardoll DM. *Nat Rev Cancer.* 2012.

# Integrative analysis reveals selective 9p24.1 amplification, increased PD-1 ligand expression, and further induction via JAK2 in nodular sclerosing Hodgkin lymphoma and primary mediastinal large B-cell lymphoma

M Green, Blood, 2010

- Chromosome 9p24.1 amplification and increased expression of PD-L1 in HL and MLBCL cell lines



# Clinical results with CPI

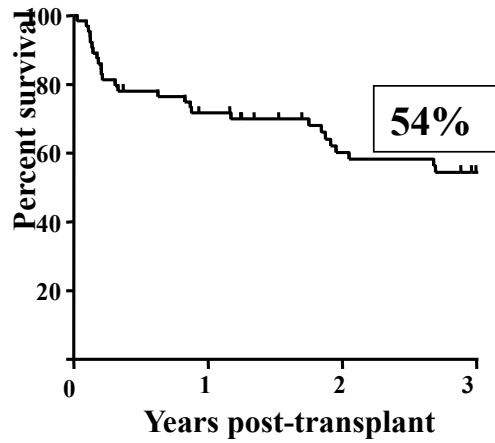
|                     | Ansell 2014   | Younes 2016 | Armand 2016 |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
|---------------------|---|-------------|-------------|--------------|----|----|----|----|----|----|----|----|----|----|-----------------|----|----|----|----|----|----|---|---|---|---|
| <b>N</b>            | 23  | 80          | 31          |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| <b>A</b>            |   |             |             |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| <b>C</b>            |   |             |             |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| <b>C</b>            | <table border="1"> <thead> <tr> <th>Time (weeks)</th> <th>0</th> <th>10</th> <th>20</th> <th>30</th> <th>40</th> <th>50</th> <th>60</th> <th>70</th> <th>80</th> <th>90</th> </tr> </thead> <tbody> <tr> <td>No. of patients</td> <td>31</td> <td>28</td> <td>23</td> <td>18</td> <td>15</td> <td>10</td> <td>7</td> <td>4</td> <td>1</td> <td>0</td> </tr> </tbody> </table> |             |             | Time (weeks) | 0  | 10 | 20 | 30 | 40 | 50 | 60 | 70 | 80 | 90 | No. of patients | 31 | 28 | 23 | 18 | 15 | 10 | 7 | 4 | 1 | 0 |
| Time (weeks)        | 0   | 10          | 20          | 30           | 40 | 50 | 60 | 70 | 80 | 90 |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| No. of patients     | 31  | 28          | 23          | 18           | 15 | 10 | 7  | 4  | 1  | 0  |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| <b>PFS</b>          | 86% 24w   | 77% 6M      | 46% 1y      |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |
| <b>Response max</b> | /   | 2.1M        | /           |              |    |    |    |    |    |    |    |    |    |    |                 |    |    |    |    |    |    |   |   |   |   |

## Haplo vs HLAid for HL relapsed post-HDC

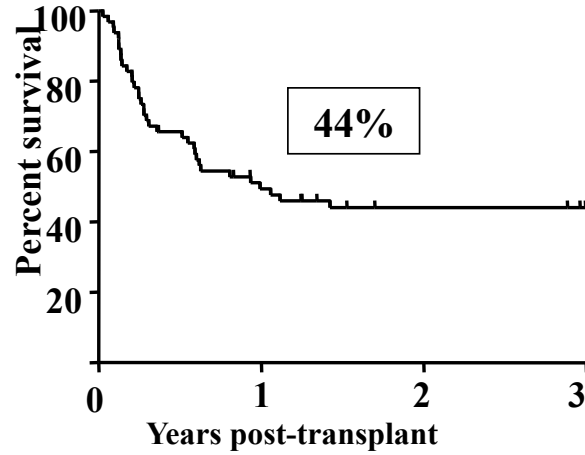
|                                | N 64           | HLAid n 34     | Haplo n 30     | p       |
|--------------------------------|----------------|----------------|----------------|---------|
| Median age                     | 31 (18-66)     | 31 (18-62)     | 31 (19-66)     | 0,8     |
| Sex M/F                        | 30/20          | 24/10          | 20/10          | 0,7     |
| N lines CT (median)            | 4 (3-13)       | 4 (3-13)       | 4 (3-8)        | 0,8     |
| Median time after Auto-Tx (ms) | 19,1 (1,7-130) | 17,6 (1,9-117) | 20,5 (1,7-130) | 0,1     |
| Disease status at ALLO         |                |                |                |         |
| CR                             | 25 (39%)       | 11 (32%)       | 14 (46%)       | 0,4     |
| PR                             | 23 (37%)       | 13 (38%)       | 10 (33%)       |         |
| SD/PD                          | 16 (24%)       | 10 (30%)       | 6 (21%)        |         |
| Stem cell source               |                |                |                |         |
| BM                             | 25 (40%)       | 2 (6%)         | 23(76%)        | <0.0001 |
| PBSC                           | 39 (60%)       | 32 (94%)       | 7 (24%)        |         |
| HCT-CI (n= 45)                 | N=49           | N= 20          | N= 29          | 0,08    |
| 0-1                            | 16 (25%)       | 3 (8%)         | 13 (43%)       |         |
| 2                              | 13 (20%)       | 6 (16%)        | 7 (23%)        |         |
| >3                             | 20 (31%)       | 11 (32%)       | 9 (30%)        |         |
| Conditioning                   |                |                |                | 0,1     |
| NMA                            | 35 (54%)       | 15(44%)        | 20 (66%)       |         |
| RIC                            | 25 (39%)       | 16 (47%)       | 9 (30%)        |         |
| MAC                            | 4 (7%)         | 3 (9%)         | 1 (4%)         |         |

# Haplo vs HLAid for HL relapsed post-HDC

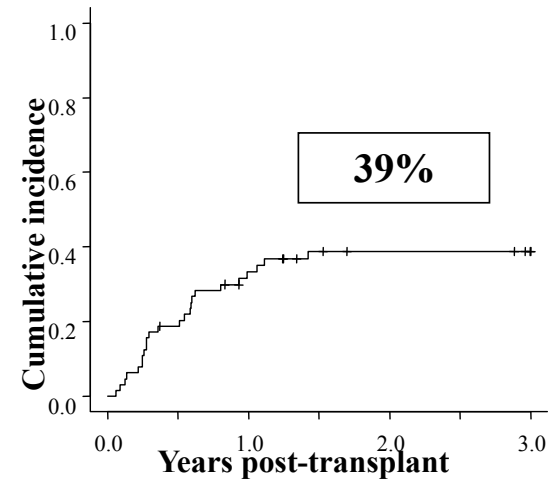
3y-OS



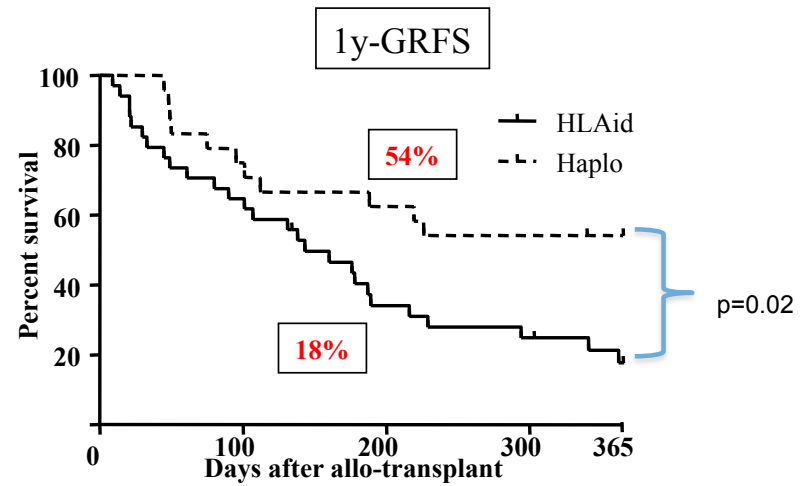
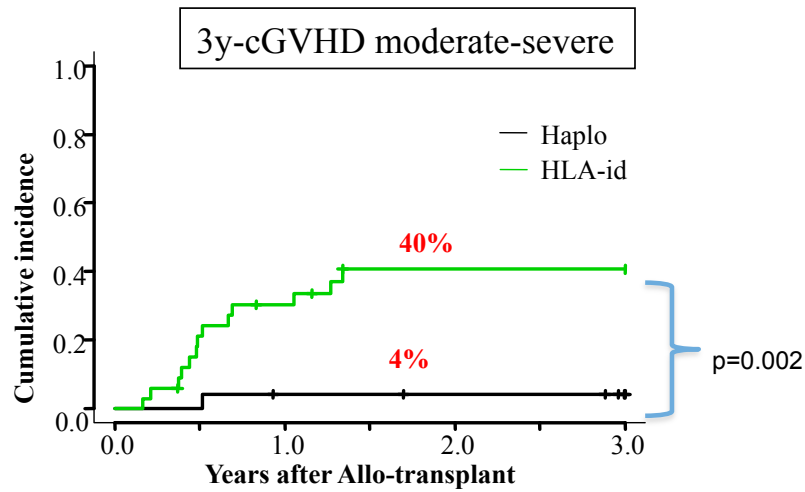
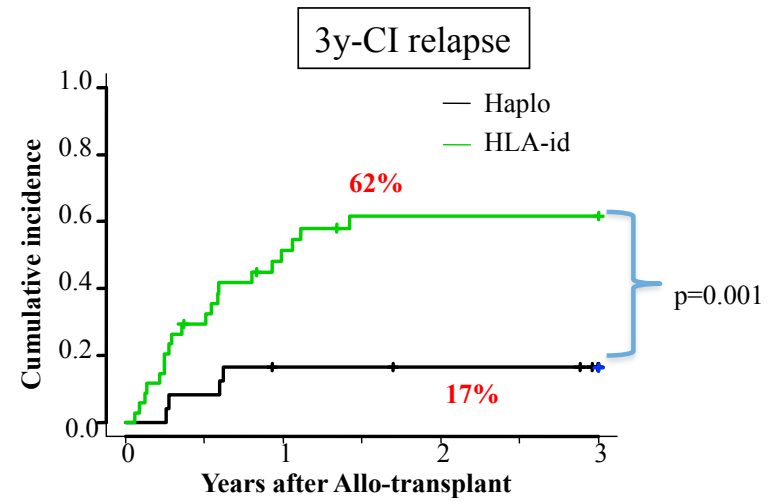
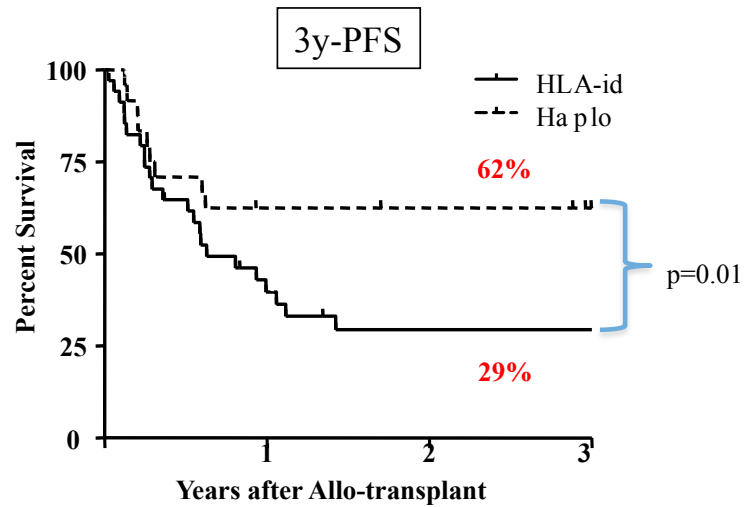
3y-PFS



3y-CI relapse

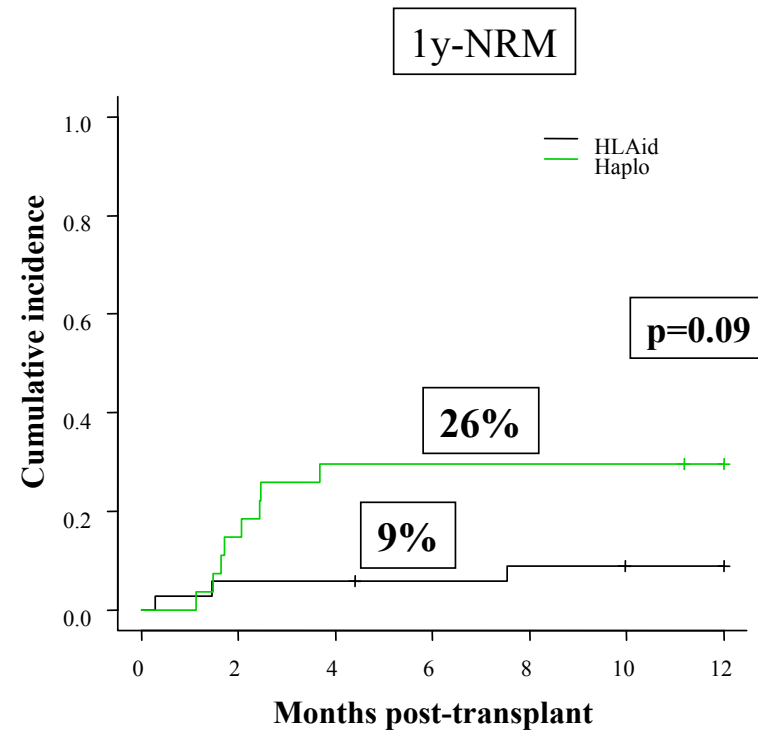
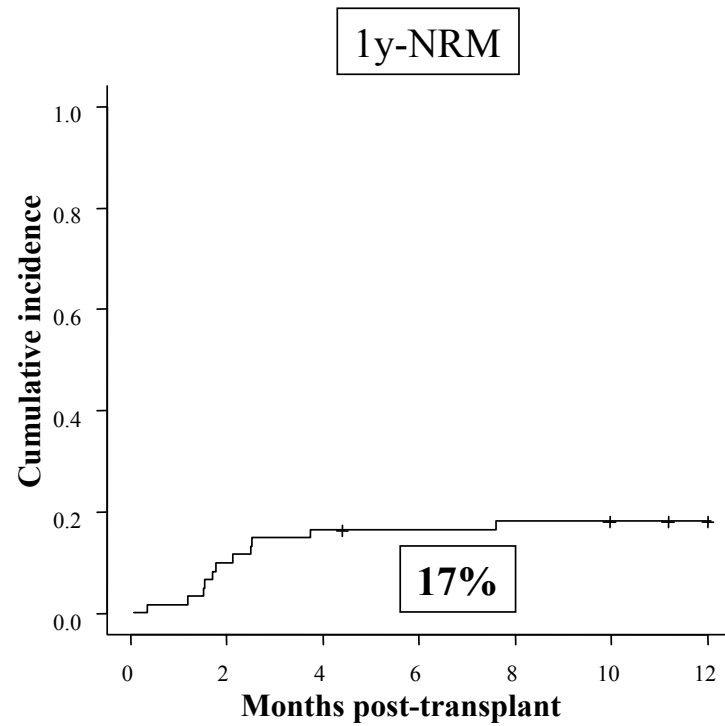


# Haplo vs HLAid for HL relapsed post-HDC





# Haplo vs HLAid for HL relapsed post-HDC



## Haplo vs HLAid for HL relapsed post-HDC

| Characteristics          | 3y-OS<br>HR          | p                 | 3y-PFS<br>HR        | p            | 1y-NRM<br>HR      | p           |
|--------------------------|----------------------|-------------------|---------------------|--------------|-------------------|-------------|
| <b>Not CR vs CR</b>      | <b>12 (3.5-41.2)</b> | <b>&lt;0.0001</b> | <b>4.6 (1.9-11)</b> | <b>0.001</b> | <b>8 (1.4-43)</b> | <b>0.01</b> |
| Haplo vs HLAid           | 1 (0.1-10.1)         | 0.9               | 0.4 (0.05-3.8)      | 0.4          | 9.4 (0.4-200)     | 0.1         |
| PBSC vs BM               | 0.4 (0.05-4)         | 0.4               | 0.4 (0.05-4.1)      | 0.4          | 2.6 (0.1-54)      | 0.5         |
| RIC/MAC vs NMA           | 2.3 (0.8-6.2)        | 0.1               | 3.2 (1.1-8.6)       | <b>0.02</b>  | 9.1 (0.9-87)      | <b>0.05</b> |
| HCT-CI $\geq 3$ vs $< 3$ | 2.7 (0.9-8.3)        | 0.06              | 1.8 (0.6-4.7)       | 0.2          | 2.8 (0.6-11.7)    | 0.1         |

|  | All pts<br>N= 198 | HLA-id<br>N= 133 | Haplo<br>N= 65 | p      |
|--|-------------------|------------------|----------------|--------|
| Age, years (median, range)               | 32 (18-66)        | 32 (18-65)       | 31 (18-65)     | 0.8    |
| Sex M/F                                  | 113/85            | 77/56            | 36/29          | 0.7    |
| Number of CT lines (median, range)       | 2 (1-12)          | 2 (1-2)          | 4 (2-12)       | <0.001 |
| Relapse after HDC                        | 170/198 (86%)     | 110/133 (83%)    | 60/65 (92%)    |        |
| <b>Disease status at transplantation</b> |                   |                  |                |        |
| CR                                       | 119 (60%)         | 82 (62%)         | 37 (60%)       |        |
| PR                                       | 79 (40%)          | 51 (38%)         | 28 (40%)       |        |
| <b>Donors</b>                            |                   |                  |                | /      |
| HLA sibling                              | /                 | 57 (43%)         | /              |        |
| MUD                                      | /                 | 76 (57%)         | /              |        |
| Haplo                                    |                   | /                | 48             |        |
| <b>Stem cell source</b>                  |                   |                  |                | <0.001 |
| PBSC                                     | 119 (60%)         | 114 (86%)        | 58 (89%)       |        |
| BM                                       | 61 (31%)          | 18 (13%)         | 43 (92%)       |        |
| miss                                     | 10 (9%)           | 1 (1%)           | /              |        |
| <b>ATG prophylaxis GVHD</b>              |                   |                  |                | /      |
| No                                       | /                 | 57               | /              |        |
| Yes                                      | 76                | 76               | /              |        |
| <b>Conditioning regimens</b>             |                   |                  |                | <0.001 |
| NMAC                                     | 58 (29%)          | /                | 58 (89%)       |        |
| RIC                                      | 101 (51%)         | 94 (71%)         | 7 (11%)        |        |
| MAC                                      | 39 (20%)          | 39 (29%)         | /              |        |

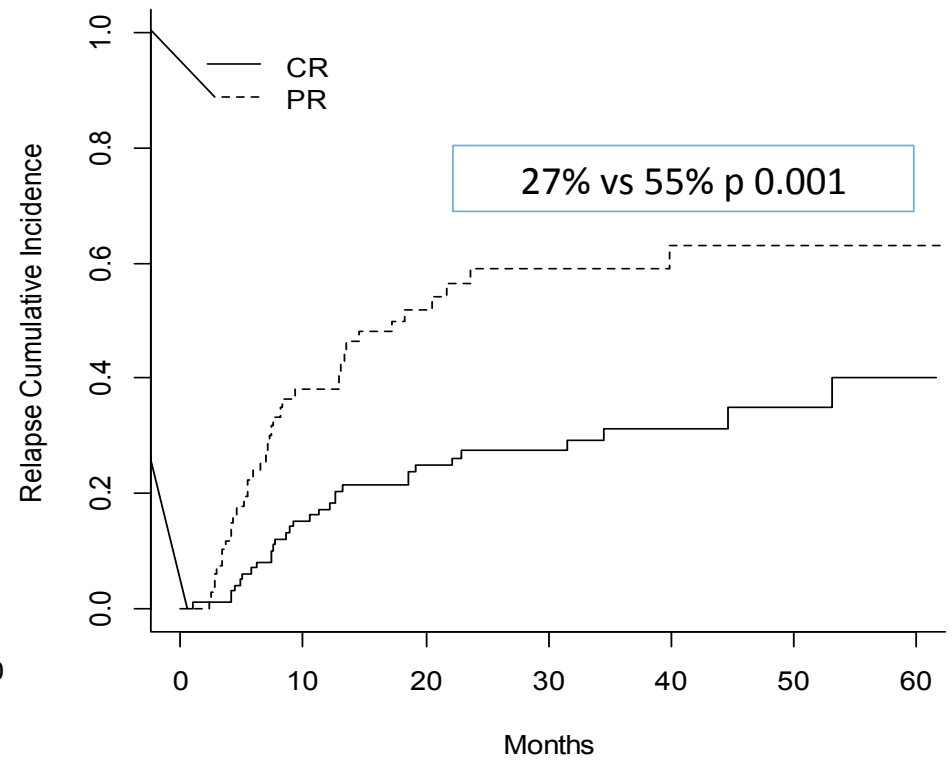
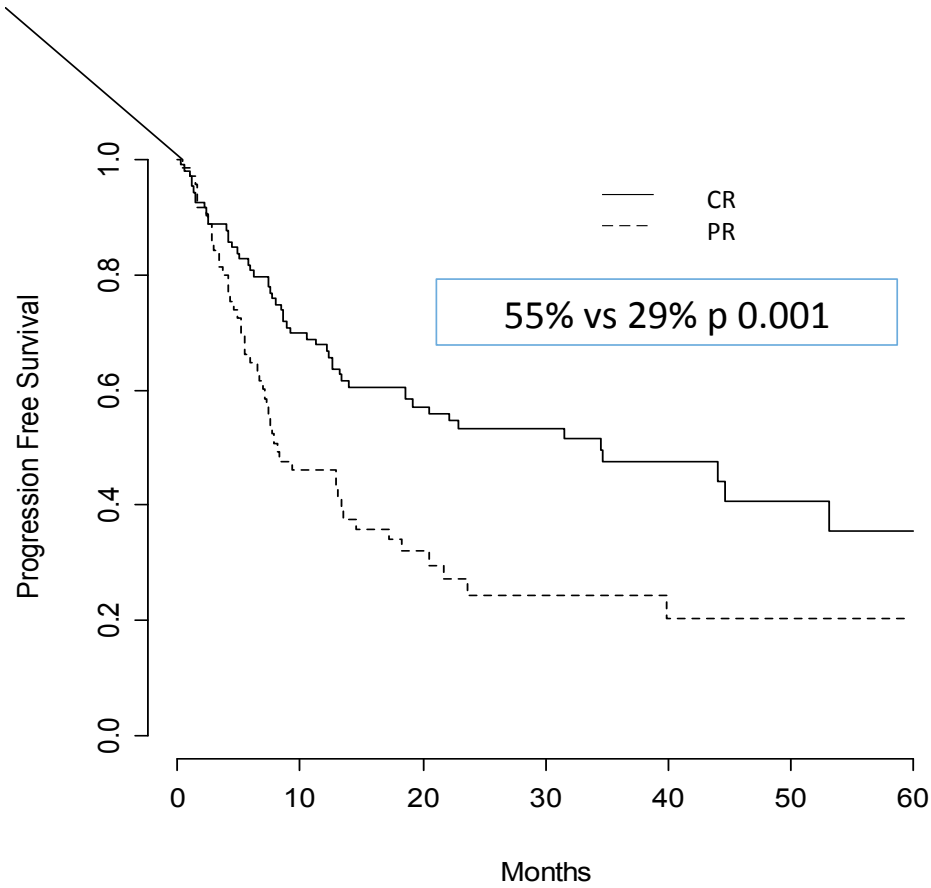
## Haplo vs HLAid CR and PR

Median follow up:  
31M (range 0.2-74.1)

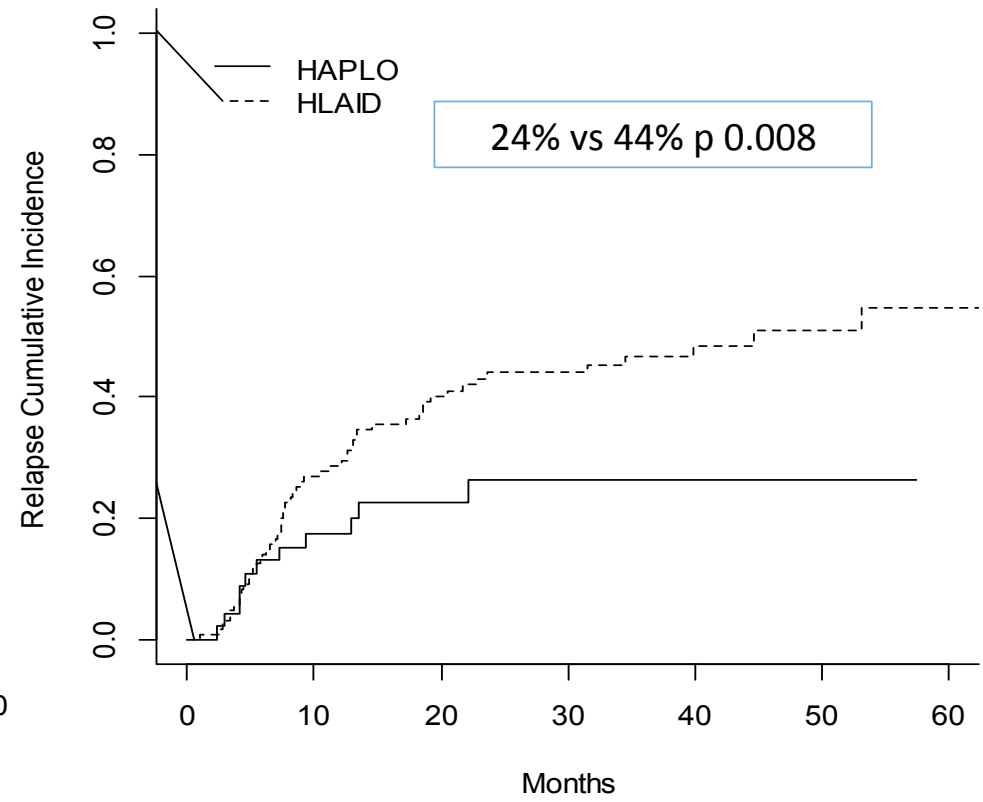
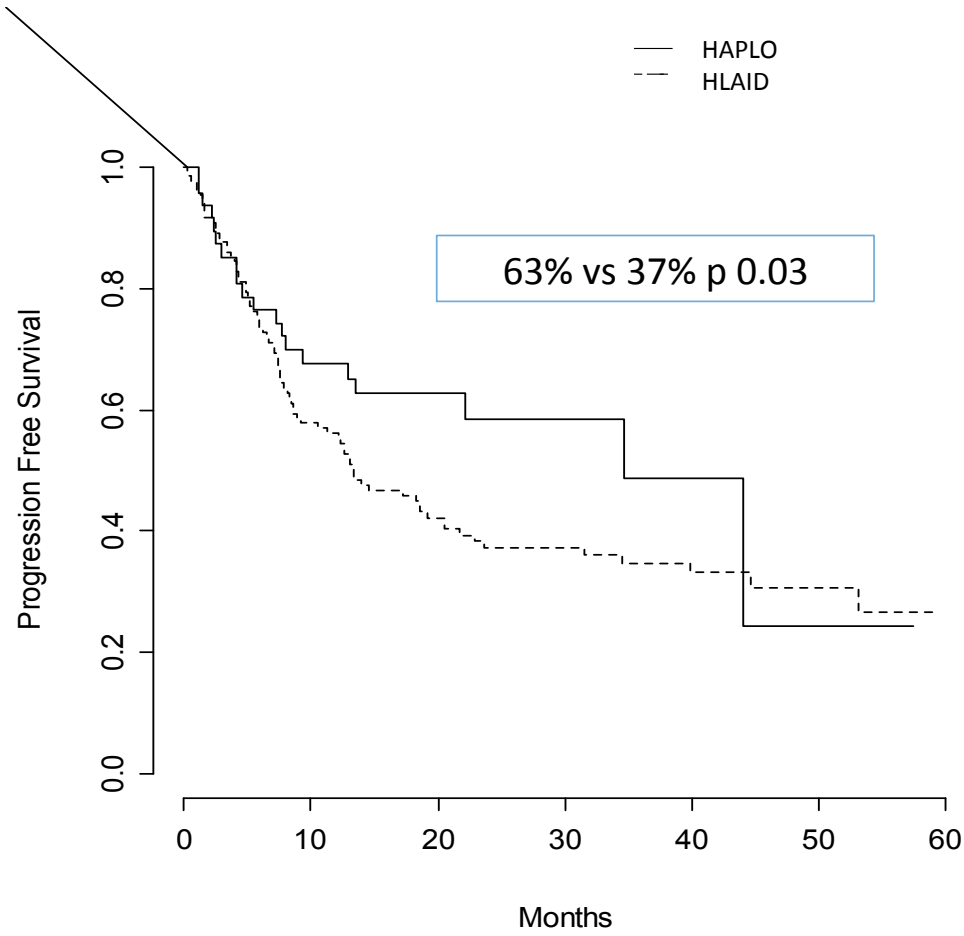
## HAPLO vs HLAid CR and PR

|   | N                    | 2y PFS                   | p                  | 2y OS                    | p           | 2y RI                    | p                 | 1y NRM                   | p   |
|---|----------------------|--------------------------|--------------------|--------------------------|-------------|--------------------------|-------------------|--------------------------|-----|
| All   | 198                  | 45%                      |                    | 66%                      |             | 38%                      |                   | 14%                      |     |
| CR vs PR  | 119 vs 79            | 55% vs 29%               | <b>0.001</b>       | 74% vs 55%               | <b>0.03</b> | 27% vs 55%               | <b>&lt; 0.001</b> | 13% vs 16%               | 0.8 |
| Haplo vs HLAid  | 65 vs 133            | 63% vs 37%               | <b>0.03</b>        | 67% vs 63%               | 0.6         | 24% vs 44%               | <b>0.008</b>      | 13% vs 15%               | 0.9 |
| CR Haplo vs<br>CR HLAid vs<br>PR Haplo vs<br>PR HLAid | 37<br>82<br>28<br>51 | 75%<br>47%<br>44%<br>22% | <b>&lt; 0.001*</b> | 83%<br>67%<br>58%<br>54% | 0.1         | 11%<br>34%<br>44%<br>60% | <b>&lt; 0.001</b> | 14%<br>13%<br>11%<br>18% | 0.8 |

# HAPLO vs HLAid CR and PR



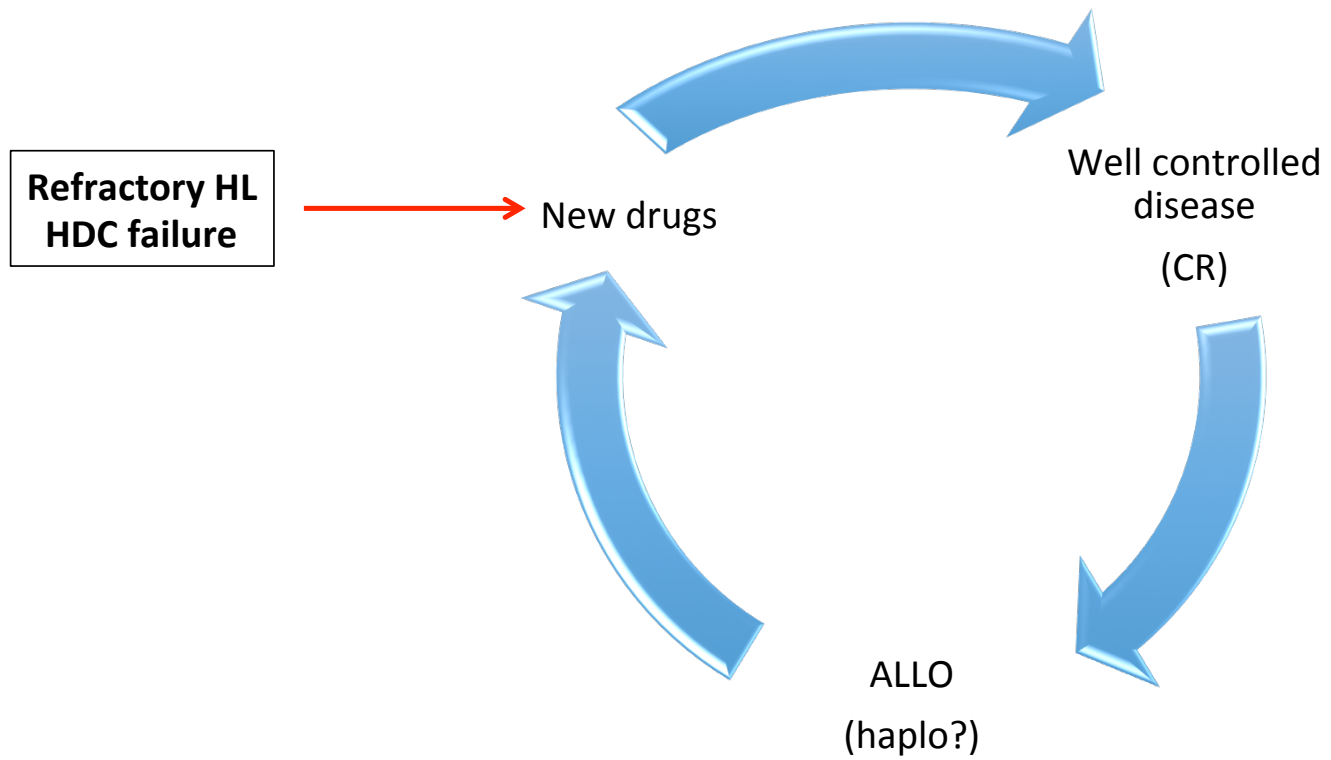
# HAPLO vs HLAid CR and PR



## HAPLO vs HLAid CR and PR

| MV                     | HR   | CI95%     | P value |
|------------------------|------|-----------|---------|
| <b>PFS</b>             |      |           |         |
| Disease Status at ALLO | 0.51 | 0.34-0.75 | <0.001  |
| Haplo vs HLAid         | 0.56 | 0.35-0.89 | 0.014   |
| <b>OS</b>              |      |           |         |
| Disease Status at ALLO | 0.57 | 0.35-0.93 | 0.023   |
| Age                    | 0.04 | 1.00-1.05 | 0.043   |
| <b>Relapse</b>         |      |           |         |
| Haplo vs HLAid         | 0.37 | 0.23-0.60 | <0.001  |
| Disease Status at ALLO | 0.43 | 0.43-0.79 | 0.006   |

# ALLO in HL





## Conclusions

- Allo-SCT must be considered for HL patients relapsed or progressed after HDC or refractory to salvage therapies
- New drugs should be integrated in the treatment strategy before allo-SCT and, if possible, after allo-SCT for those at higher risk of relapse
- Allo-SCT toxicity can be modified by immunomodulatory drugs, but this cannot be an exclusion criterion.
- Similarly, immunomodulatory drugs used after allo-SCT, can be induce important side effects.