

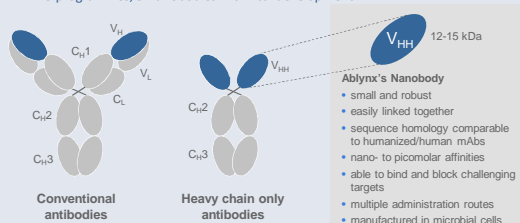


An innovative approach for detecting neutralizing antibodies directed to antibody-derived therapeutics based on the conventional bridging anti-drug antibody (ADA) assay format

Veerle Snoeck, Carlo Boutton, Annelies Coddens, Benedicte Brackeva, Robbe D'Hondt, Gregory Daelman, Tom F. Verhaeghe, Andreia Correia, Hans Ulrichts, Peter Casteels, Marie-Ange Buyse
Ablynx NV, Zwijnaarde, Belgium

Background: Nanobodies

- The isolated variable domains (VHH) of camelid heavy-chain only antibodies are stable and fully functional
- Nanobodies represent the next generation of antibody-derived biologics
- > 45 programmes, 8 Nanobodies in clinical development



Alternative NAB assay – fit for purpose using mAbs

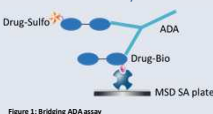
Fit for purpose of assay format demonstrated using a panel of established neutralizing and non-neutralizing Ab

- Neutralizing potential of the monoclonal antibodies (mAb) is based on the ability to block target interaction as determined in the CLBA (in buffer without pre-treatment)
- NAB are detected at the same sensitivity as compared to the ADA assay
- Non-neutralizing Ab are left undetected; some residual binding can be detected at very high Ab concentrations, however these levels are not expected to be clinically relevant

| Neutralizing Ab | Assay | Assay response (ECL) at mAb concentration (ng/mL) | | | | | | | | Sensitivity | Sensitivity in conventional NAB assay |
|----------------------------|-------|---|-------|------|------|------|-----|-----|-----|-------------|---------------------------------------|
| | | 20000 | 4000 | 1000 | 500 | 250 | 125 | 63 | 0 | | |
| mAb 1 | ADA | 19714 | 4122 | 1140 | 599 | 375 | 242 | 178 | 109 | < 63 ng/mL | |
| | NAB | 23151 | 4623 | 1266 | 640 | 383 | 246 | 173 | 92 | < 63 ng/mL | 556 ng/mL |
| mAb 2 | ADA | 35882 | 10405 | 2796 | 1405 | 784 | 468 | 273 | 156 | < 63 ng/mL | |
| | NAB | 62575 | 12217 | 3097 | 1596 | 854 | 466 | 278 | 89 | < 63 ng/mL | 556 ng/mL |
| mAb 3 | ADA | 646 | 214 | 139 | 102 | 118 | 117 | 110 | 116 | 400 ng/mL | |
| | NAB | 671 | 203 | 127 | 105 | 103 | 99 | 94 | 94 | 543 ng/mL | > 5 µg/mL |
| mAb 4 | ADA | 757 | 250 | 147 | 136 | 125 | 119 | 116 | 107 | 165 ng/mL | |
| | NAB | 807 | 251 | 140 | 127 | 117 | 108 | 103 | 95 | 111 ng/mL | > 5 µg/mL |
| Non-neutralizing Ab | | | | | | | | | | | |
| mAb 5 | ADA | 14731 | 5182 | 1811 | 935 | 530 | 301 | 209 | 102 | < 63 ng/mL | |
| | NAB | 128 | 101 | 94 | 88 | 88 | 89 | 92 | 96 | 889 ng/mL | N/A |
| mAb 6 | ADA | 1624 | 635 | 296 | 165 | 138 | 129 | 133 | 120 | < 63 ng/mL | |
| | NAB | 96 | 92 | 95 | 93 | 96 | 97 | 99 | 97 | > 20 µg/mL | N/A |
| mAb 7 | ADA | 1197 | 344 | 170 | 141 | 138 | 116 | 120 | 113 | 211 ng/mL | |
| | NAB | 96 | 89 | 80 | 83 | 84 | 83 | 84 | 85 | > 20 µg/mL | N/A |
| mAb 8 | ADA | 10620 | 20679 | 5109 | 2625 | 1314 | 733 | 446 | 110 | < 63 ng/mL | |
| | NAB | 117 | 92 | 83 | 80 | 82 | 84 | 89 | 89 | 11023 ng/mL | N/A |

Background: Immunogenicity testing

- Neutralizing antibodies (NAB) are defined as antibodies (Ab) able to affect the function of the Nanobodies by blocking its ability to bind to target, including Ab binding to complementary determining region (CDR)
- Development of a NAB assay based on drug-target interaction is challenging and often requires extensive pre-treatment steps in order to obtain the required drug and target tolerance for proper detection of NAB during clinical trials
- Such an assay set-up often introduces a sensitivity difference between the ADA assay and the NAB assay
- The sensitivity gap complicates ADA data interpretation as a discrepancy between ADA and NAB results can either reflect presence of non-neutralizing antibodies only or reflect neutralizing antibodies that are left undetected in the NAB assay
- An alternative NAB assay format was developed based on the conventional bridging ADA assay format which allows unambiguous comparison of the levels of total ADA and NAB



Alternative NAB assay – fit for purpose using polyclonals

Neutralizing antibody fraction sensitively detected in polyclonal antibody samples

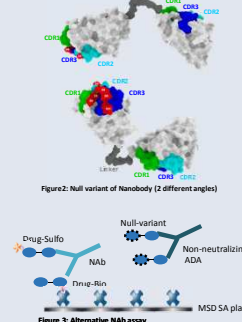
- The polyclonal rabbit Ab was specifically generated by immunization to be used as NAB positive control antibody
- The pre-clinical study samples (rhesus monkey) originating from a disease model shown to be prone to development of ADA, were shown to contain neutralizing activity via PD and efficacy markers
- The neutralizing fraction within the ADA positive samples (Rabbit pAb and pre-clinical study samples) can be determined by titer and/or sensitivity determination

| Rabbit pAb | Assay | Assay response (ECL) at pAb concentration (ng/mL) | | | | | | | | Sensitivity | Sensitivity in conventional NAB assay |
|------------------------------------|-------|---|-------|------|------|-------|--------|--------|---------|-------------|---------------------------------------|
| | | 4000 | 1000 | 500 | 250 | 125 | 63 | 0 | | | |
| ADA | ADA | 4694 | 1192 | 659 | 230 | 135 | 117 | 114 | 105 | < 5.0 ng/mL | |
| NAB | NAB | 1520 | 416 | 267 | 129 | 98 | 88 | 95 | 92 | 31 ng/mL | 5 µg/mL |
| Rhesus Monkey study samples | | | | | | | | | | | |
| Sample 1 | Assay | Assay response (ECL) at pAb dilution | | | | | | | | Sensitivity | LOD (CFU) |
| | | 100 | 400 | 1600 | 6400 | 25600 | 102400 | 409600 | 1638400 | | |
| ADA | ADA | 253499 | 25133 | 4526 | 812 | 236 | 126 | 105 | 5.1 | | |
| NAB | NAB | 95676 | 15281 | 3032 | 581 | 176 | 106 | 92 | 5.0 | | |
| ADA | ADA | 714810 | 61990 | 9795 | 1711 | 389 | 158 | 110 | 5.5 | | |
| NAB | NAB | 268224 | 41922 | 7437 | 1363 | 309 | 123 | 99 | 5.4 | | |
| ADA | ADA | 81076 | 8794 | 1696 | 333 | 144 | 112 | 98 | 4.9 | | |
| NAB | NAB | 35866 | 6523 | 1197 | 274 | 125 | 96 | 87 | 4.8 | | |

The alternative NAB assay

The proposed alternative NAB assay is based on the conventional bridging ADA assay format

- An excess amount of the null variant of the Nanobody, i.e. a variant of the Nanobody which is non-functional for target binding and which is identical to the Nanobody with the exception of altered CDR's of the target binding Nanobody domain, is added to the reagent master mix
- Non-neutralizing antibodies are complexed with the null variant of the Nanobody and are left undetected (similar to a drug-displacement set-up as confirmatory assay of the conventional ADA assay)
- Positive assay signals reflect antibodies with neutralizing potential only



ADA and alternative NAB assay qualification data

Assay qualification reveals similar precision, sensitivity and drug tolerance characteristics as compared to the ADA assay

- Similar sensitivity and drug tolerance, compliant to current regulatory guidelines: < 100 ng/mL positive control in presence of highest anticipated drug levels

| Conc. mAb 2 (ng/mL) | ADA - Mean responses (ECL) | | | | | Alternative NAB - Mean responses (ECL) | | | | |
|---------------------|----------------------------|--------|-----------|--------|---------|--|--------|-----------|--------|---------|
| | 30.0 | 15.0 | 0.6 µg/mL | 0.03 | No drug | 30.0 | 15.0 | 0.6 µg/mL | 0.03 | No drug |
| 10000.0 | 27266 | 23744 | 41832 | 38100 | 34203 | 32166 | 38126 | 44809 | 67705 | 64243 |
| 5000.0 | 7264 | 9463 | 10162 | 10956 | 12297 | 7329 | 8712 | 13118 | 16188 | 16898 |
| 2000.0 | 761 | 881 | 858 | 1210 | 1470 | 858 | 966 | 1164 | 1649 | 1747 |
| 1000.0 | 441 | 525 | 512 | 545 | 745 | 456 | 482 | 619 | 817 | 1044 |
| 500.0 | 185 | 220 | 228 | 207 | 316 | 197 | 205 | 250 | 268 | 346 |
| 250.0 | 139 | 148 | 145 | 146 | 180 | 137 | 147 | 160 | 152 | 212 |
| 125.0 | 94 | 99 | 94 | 98 | 103 | 94 | 98 | 96 | 92 | 88 |
| 62.5 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 | < 36.0 |

- Target tolerance characteristics similar as the ADA assay. In case of a monomeric target, this NAB format is target tolerant
- Intra-run and inter-assay precision: ≤ 20%

| Conc. mAb 2 (ng/mL) | ADA | | | Alternative NAB | | |
|---------------------|--------------------|---------------------------------|---------------------------|--------------------|---------------------------------|---------------------------|
| | Overall mean (ECL) | Overall intra-run precision (%) | Inter-batch precision (%) | Overall mean (ECL) | Overall intra-run precision (%) | Inter-batch precision (%) |
| 20000.0 | 45942 | 4.8 | 15.8 | 64390 | 8.9 | 19.2 |
| 72.0 | 245 | 3.4 | 9.5 | 306 | 5.6 | 14.9 |
| 36.0 | 158 | 3.0 | 6.8 | 196 | 4.9 | 10.6 |
| 18.0 | 97 | 3.8 | N/A | 85 | 4.2 | N/A |

The conventional NAB assay

- Assay was developed to be fit-for-purpose according to following requirements
- Drug tolerance: predicted maximum drug levels of 30 µg/mL
- Target tolerance: predicted maximum target levels of 1500 ng/mL
- Sensitivity: 250 – 500 ng/mL positive control antibody in absence of drug

- Assay format
- Competitive ligand binding assay (CLBA) with a complex pre-treatment step to achieve drug and target tolerance



- Sensitivity gap between ADA assay and conventional NAB assay
- Both assays have been fully validated according to FDA guidelines

| (ng/mL) | NAB assay sensitivity (ng/mL) | ADA assay sensitivity (ng/mL) | Sensitivity difference ADA versus NAB assay |
|-----------|-------------------------------|-------------------------------|---|
| 60 µg/mL | 1890 | < 36 | At least 50-fold |
| 30 µg/mL | 1374 | < 36 | At least 40-fold |
| 15 µg/mL | 779 | < 36 | At least 20-fold |
| 600 ng/mL | 417 | < 36 | At least 10-fold |
| 30 ng/mL | 428 | < 36 | At least 10-fold |
| 0 | 521 | < 36 | At least 10-fold |

Conclusion

- The alternative NAB assay format is based on the conventional bridging ADA assay format. The assay is performed in presence of excess of a null variant of therapeutic drug added to the reagent master mix
- The null variant of the Nanobody is non-functional for target binding and is identical to the Nanobody with the exception of altered CDR's of the target binding Nanobody domain
- Fit for purpose of the assay format was demonstrated using an established panel of neutralizing and non-neutralizing Ab for which the (non)neutralizing potential is based on their (in)ability to block target interaction via a CLBA
- Signals generated in this assay reflect antibodies that bind to the CDR region and have a potential for neutralization.
- Non-neutralizing are left undetected as complexed with the null variant, in a way comparable to the conventional drug displacement set-up (confirmatory assay)
- Allows unambiguous immunogenicity data interpretation as potential NAB are detected with same assay characteristics as ADA assay
- While these assays represent a significant advance in methodology, the clinical relevance of the results must be viewed in relation to effects on drug PK, PD, efficacy and safety