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Preclinical evaluation of radiolabeled proteins

Anzhelika Vorobyeva, PhD, Docent

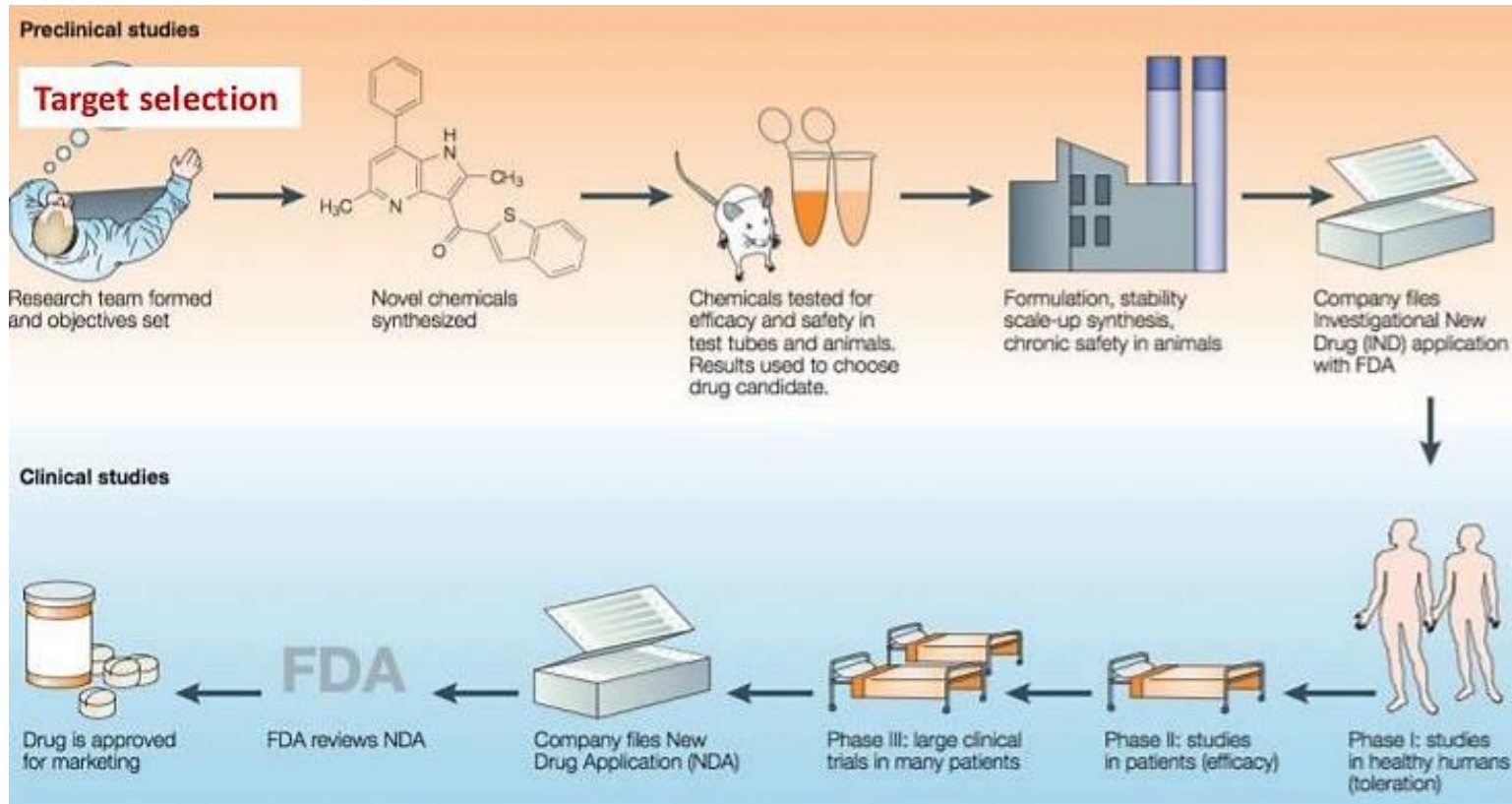
Anna Orlova, Professor

Vladimir Tolmachev, Professor

**IGP Department, Uppsala University,
Uppsala, Sweden**



Drug discovery process

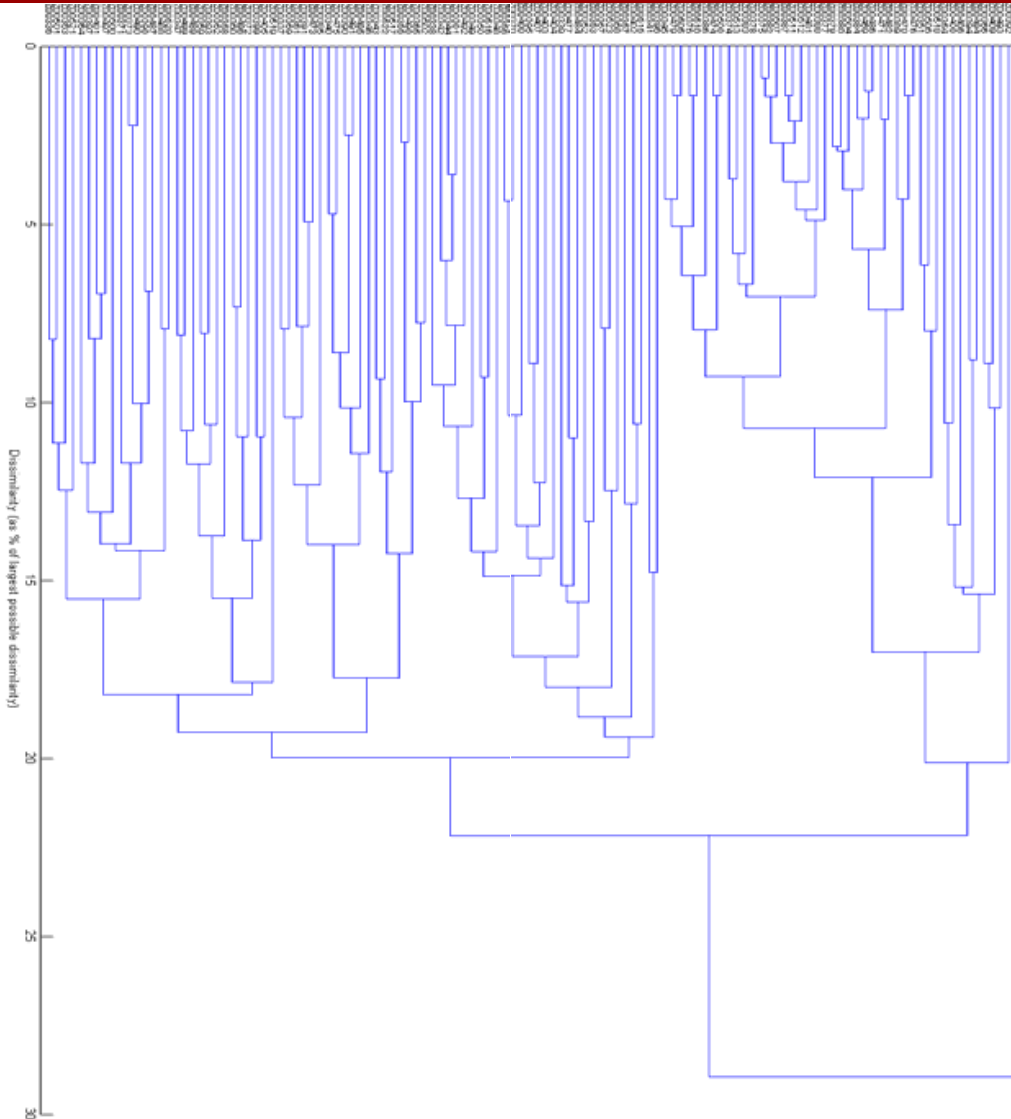


Food and Drug Administration (FDA) in the United States (or comparable agency in other countries)

The development of a new drug can take several decades and billion dollars invested



Selection of the best candidate

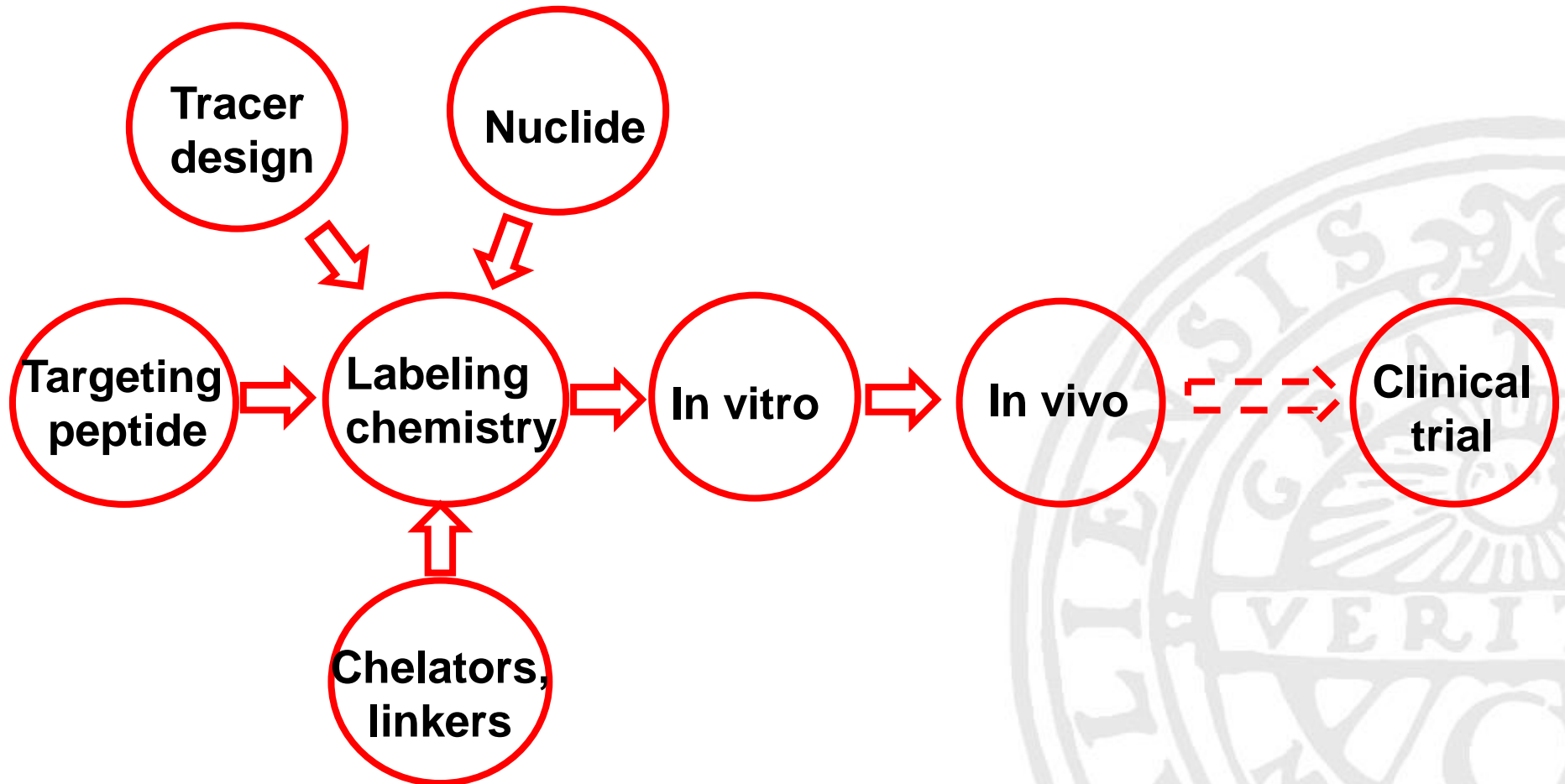


Possible selection criteria:

- Affinity
- Thermal stability
- Production
- Labeling radionuclide
- Labeling method
- Label stability
- Protein/target interaction
- Biodistribution profile
- In vivo targeting properties



Pre-clinical evaluation of radiolabeled proteins





Applications of proteins

- **Basic research**
- **Preclinical development:** pharmacokinetic studies, targeting properties, therapeutic dose
- **Clinical diagnostic:** patient stratification, therapy response
- **Treatment planning:** dosimetry estimation

What is the goal? Imaging or therapy?



Preclinical evaluation of the targeting molecule

- **Labeling of targeting molecule:** optimisation of labeling yield, confirmation of identity of labeled compound, label stability (shelf-life, stability in blood-plasma)
- **In vitro characterisation:** specificity, binding properties, cellular processing, therapy effect
- **In vivo characterisation:** normal biodistribution, target specificity, biodistribution in tumor bearing mice, confirmatorial image, dosimetry, therapy



Labeling of targeting molecule

Optimization of labeling:

Simple procedure

High yield

Reproducibility

Identity (HPLC analysis, SDS-PAGE)

Stability of labeling:

Shelf-life (stability in solution over time)

Stability under challenge (KI, histidine, EDTA)

Stability in blood plasma



Labeling of targeting molecule

- **Stability of tracer in blood**
transchelation, free radionuclide, colloids
- **Chemical modifications**
conformation, target recognition, molecular charge, lipophilicity
- **Cellular processing by malignant cells**
residualising properties of radiocatabolites
- **Retention in healthy tissues**
residualising properties of radiocatabolites

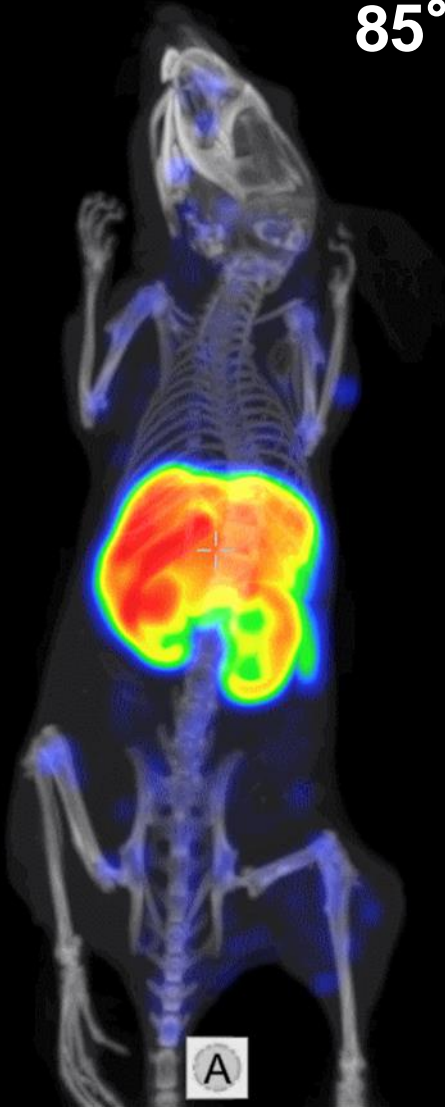


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Optimal labeling conditions!

CT+NM
Ser# 4

85° C



A

Th 0,25
Sp 0,25/0,25
Cols 140
Rows 140

Active: CT
group: none

60° C



L P

Active: CT

$^{57}\text{Co-DOTA-ZHER1}$

Garousi, unpublished data



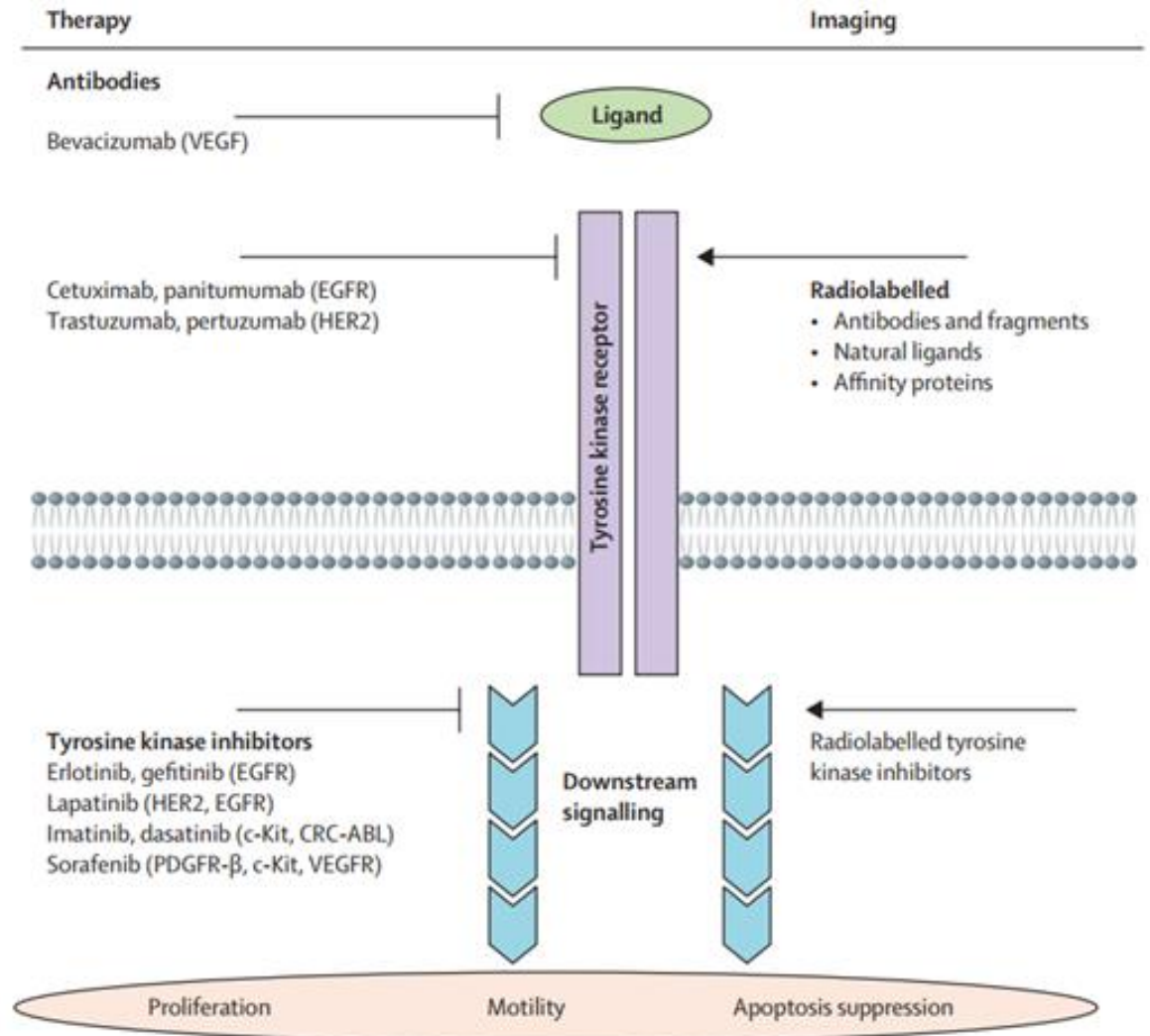
Targeting cell-surface receptors

**Example:
Receptor tyrosine kinases**

Signalling

- cell proliferation
- suppression of apoptosis
- increased motility
- recruitment of neovasculature

RTKs are targets for anticancer drugs





In vitro characterisation

Specificity:

Saturable targeting

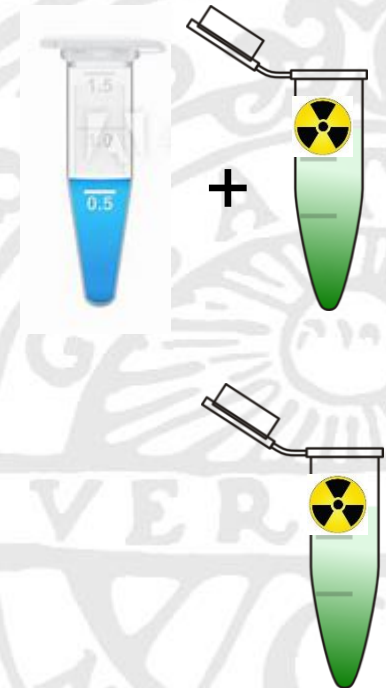
No binding to other targets



Cells expressing a target

Saturation by "cold" +
Radiolabeled compound

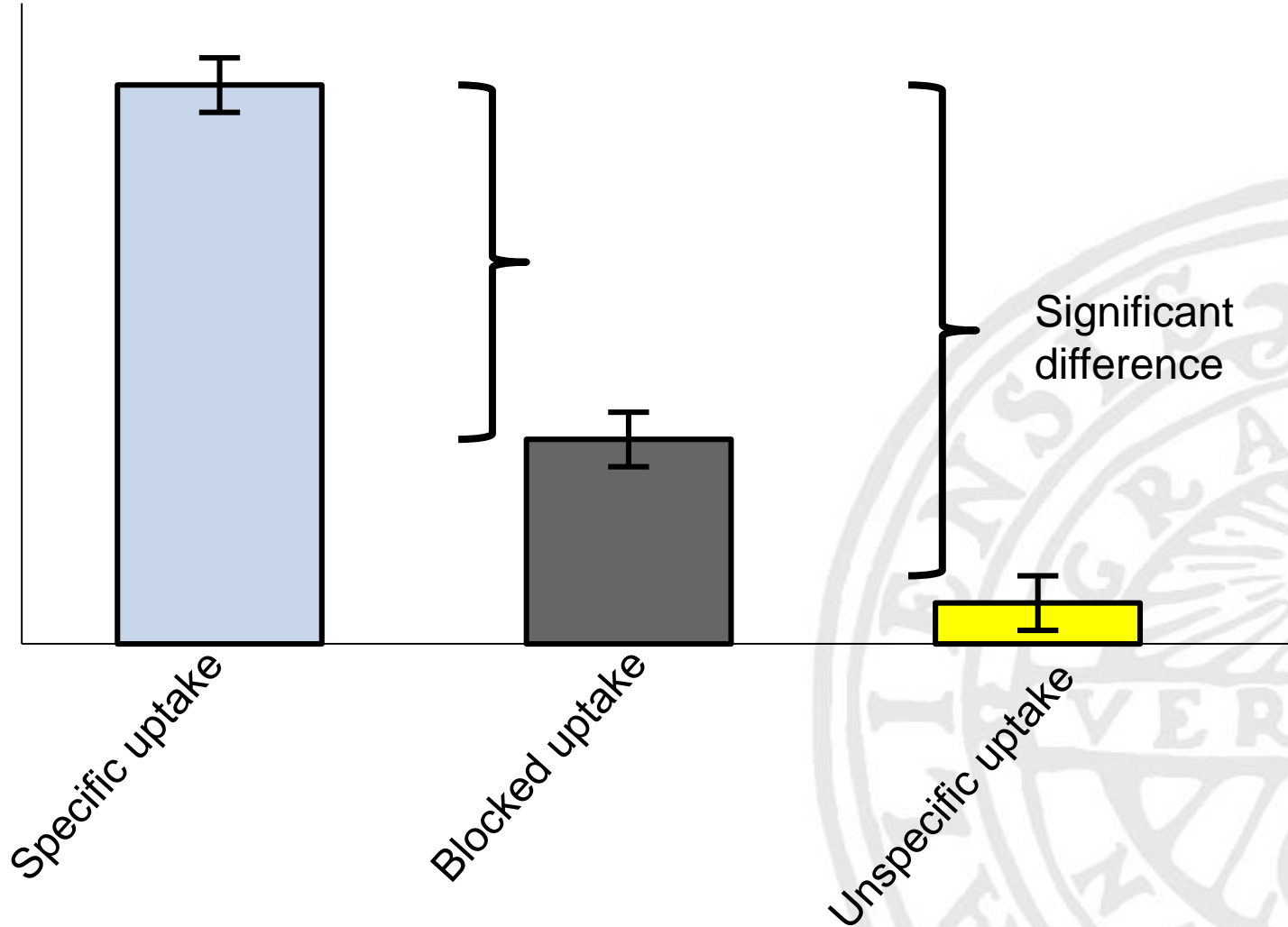
Radiolabeled compound only





In vitro specificity

Relative uptake of labeled compound





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In vitro characterisation

Specificity

Binding properties

Dissociation constant

Inhibition

Binding competent fraction

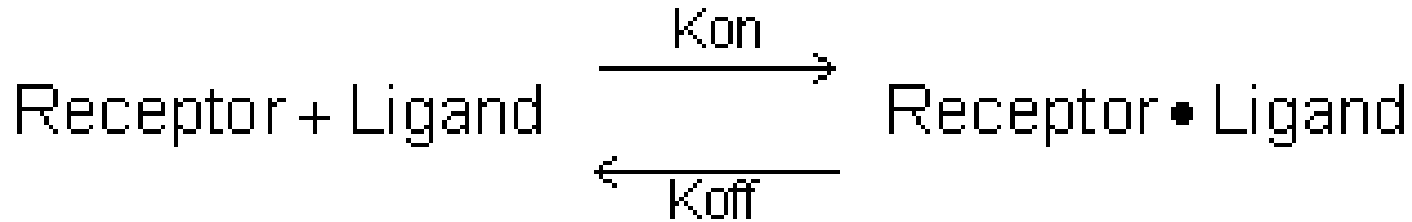
Cellular processing

Therapy effect





Equilibrium binding studies



Association of a ligand to its receptor and the dissociation of the resulting ligand-receptor complex are reversible processes that occur concomitantly until equilibrium is reached

$$\frac{[\text{Ligand}] \cdot [\text{Receptor}]}{[\text{Ligand} \cdot \text{Receptor}]} = \frac{k_{\text{off}}}{k_{\text{on}}} = K_d$$

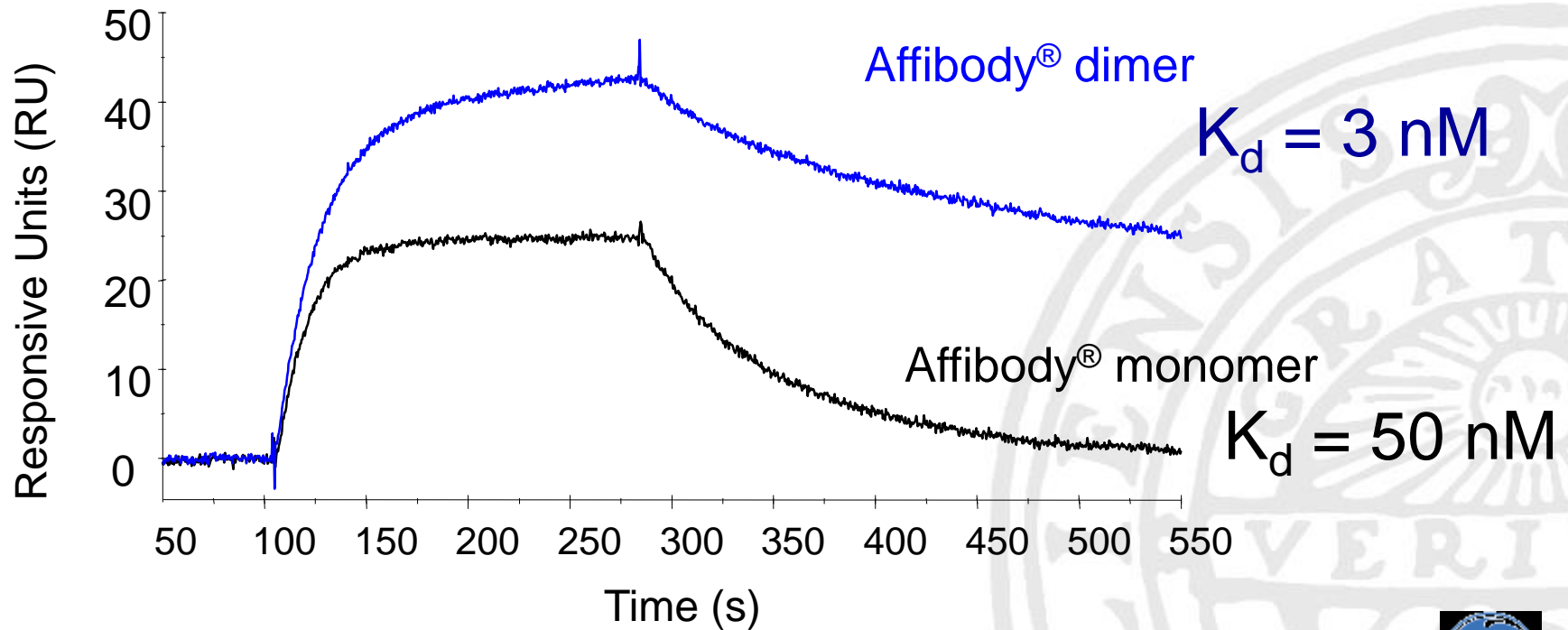
k_{on}	Association rate constant or on-rate constant
k_{off}	Dissociation rate constant or off-rate constant
K_d	Equilibrium dissociation constant



Binding properties:

Dissociation constant, K_D

BiaCore technology (receptors attached on a chip)

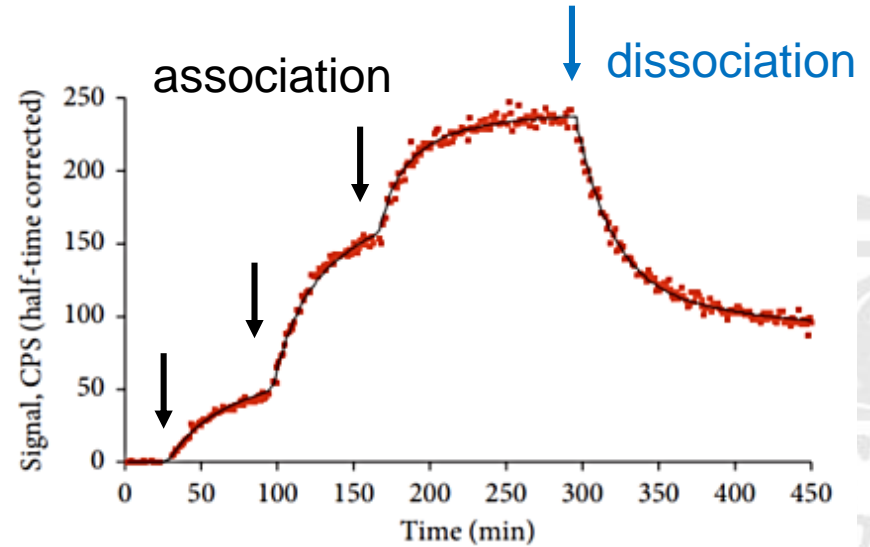
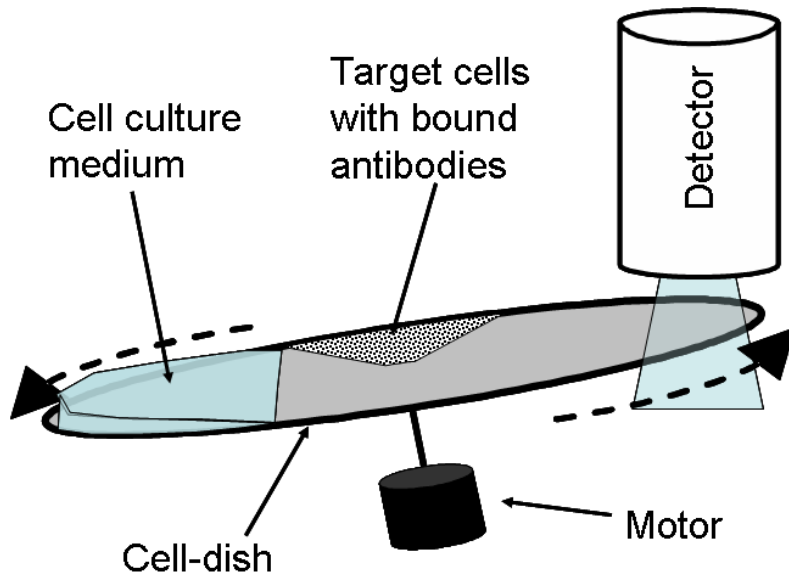




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Binding properties:

Dissociation constant, K_D



— Fitted curve
• Measured data

$$K_d = \frac{k_{\text{off}}}{k_{\text{on}}}$$

ridgeview
instruments ab

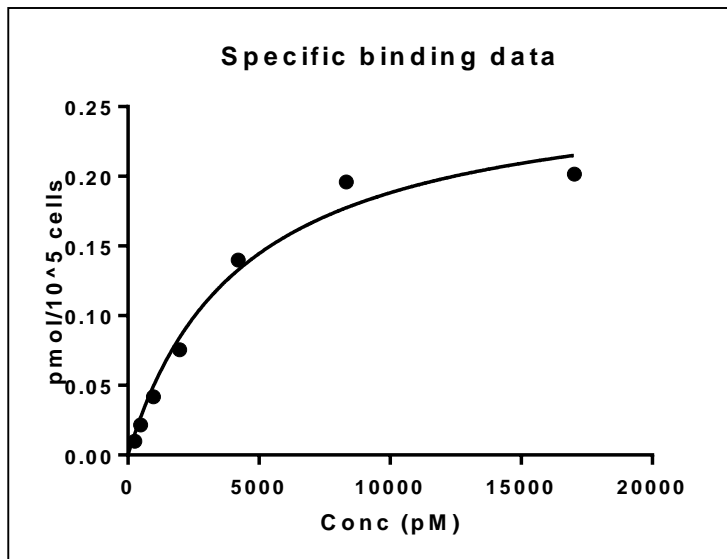


Saturation experiment:

Alternative way to measure K_d and B_{max}

1) To determine the affinity of a radioligand for a receptor

2) The density (B_{max}) of a specific receptor or receptor subtype on cells / tissues



Results of the saturation experiment are plotted with pmol/ 100 k cells on the Y axis and concentration of radioactive ligand (pM) on the X axis.

The resulting graph is a hyperbola and is called a saturation curve



Saturation experiment:

Alternative way to measure K_D and B_{max}



- Use several concentrations (up to 10) of radiolabeled ligand

$0.1 \times K_D$ ← K_D → $10 \times K_D$

- Long incubation (4-8 hours)

- For each concentration 4 dishes (3 non-blocked, 1 blocked) + 1 standard

3 dishes Nonblocked



1 dish blocked



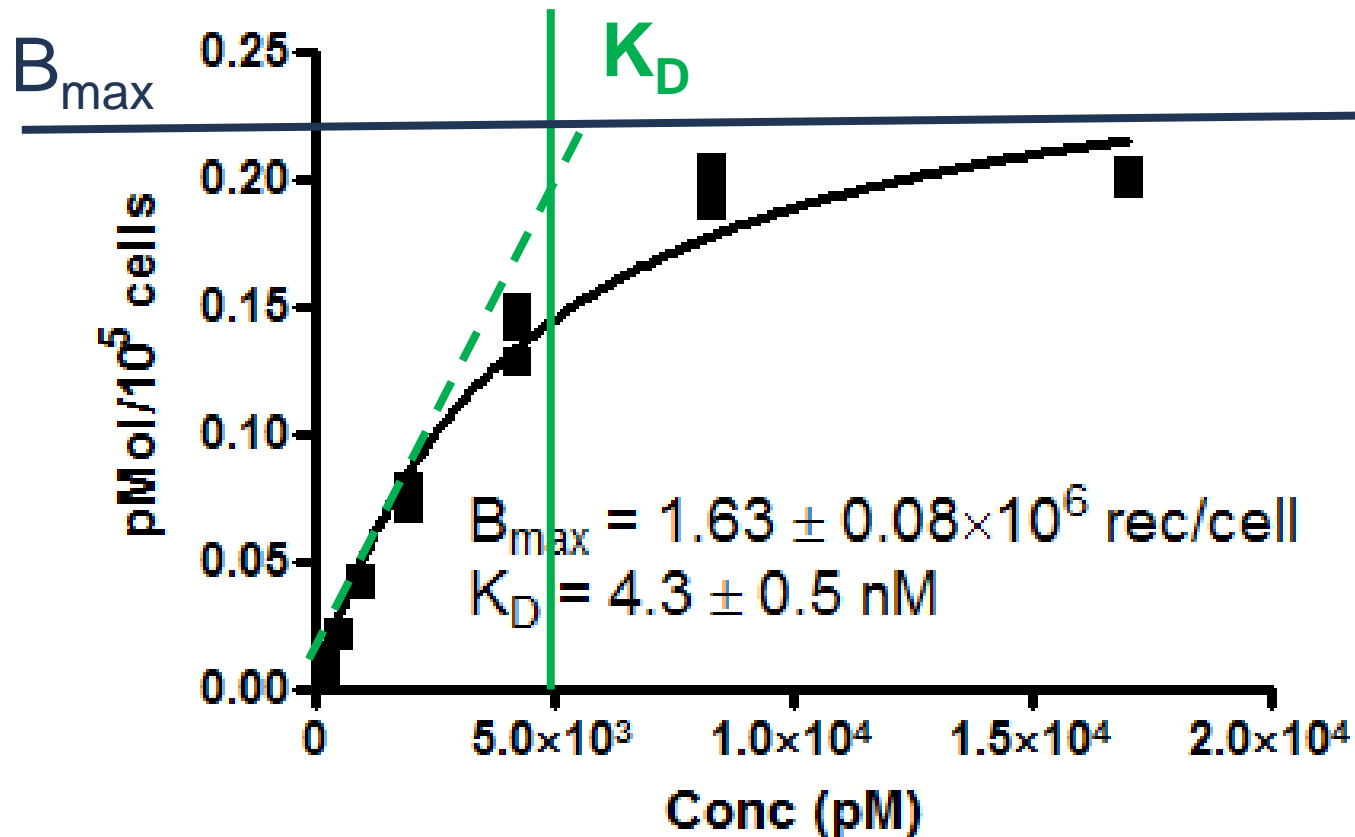


Binding properties:

Dissociation constant, K_D

Saturation analysis

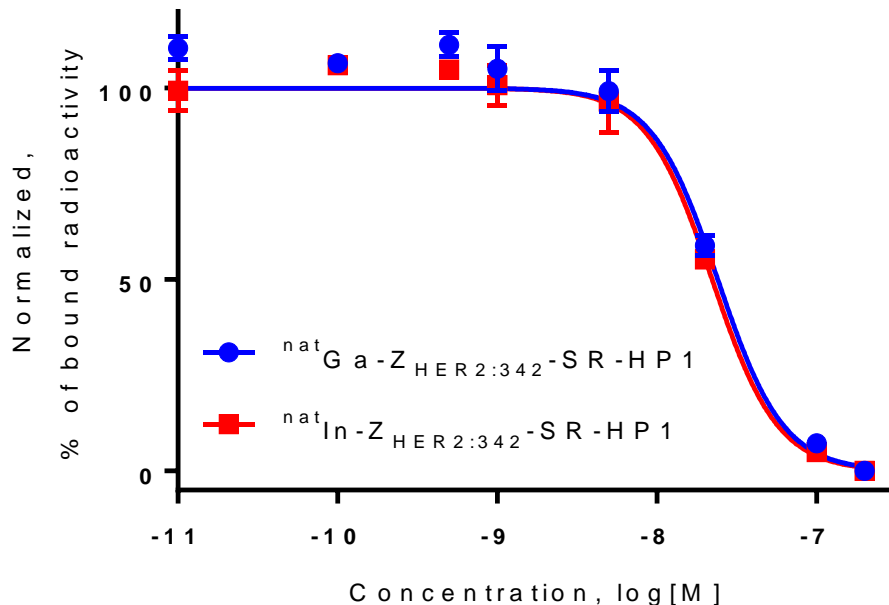
Increasing concentration of **labeled** targeting molecule





Competitive binding experiments

measure equilibrium binding of a single concentration of radioligand at various concentrations of an unlabeled competitor



- Single concentration of ^{111}In -labeled affibody
- Range of concentrations of "cold Ga" labeled affibody analogue
- Range of concentrations of "cold In" labeled affibody analogue

$$\text{IC}_{50}^{\text{nat Ga-ZHP1}} = 24.1 \pm 2.0 \text{ nM}$$

$$\text{IC}_{50}^{\text{nat In-ZHP1}} = 22.2 \pm 1.1 \text{ nM}$$

To compare the affinities of several compounds in one assay



Determination of the immunoreactive fraction of radiolabeled mAbs by linear extrapolation to binding at infinite antigen excess

- Lindmo et al. J Immunol Methods. 1984 Aug 3;72(1):77-89.

Lindmo assay

- For properly performed conjugation procedures, immunoreactive fractions of about 0.9 were obtained, but a **prolonged chloramine-T reaction for ^{125}I -labeling resulted in an immunoreactive fraction of 0.6**
- Due to its principle of determining binding at infinite antigen excess, the present method is insensitive to variation in the actual amounts of cells and antibody used, as well as the incubation time. We therefore recommend it as a **quality control procedure for radiolabeled antibodies**

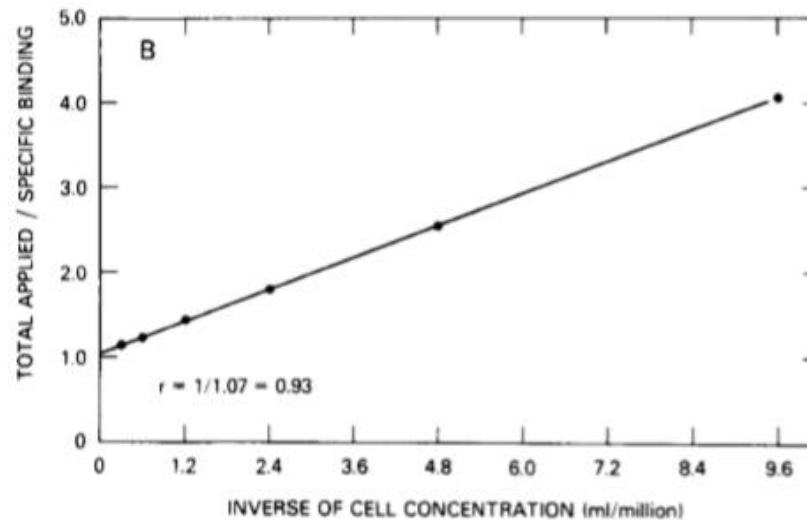
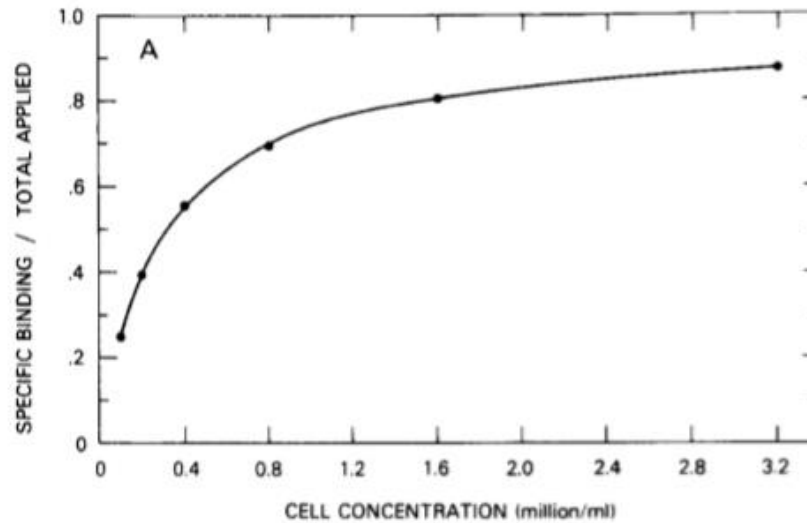


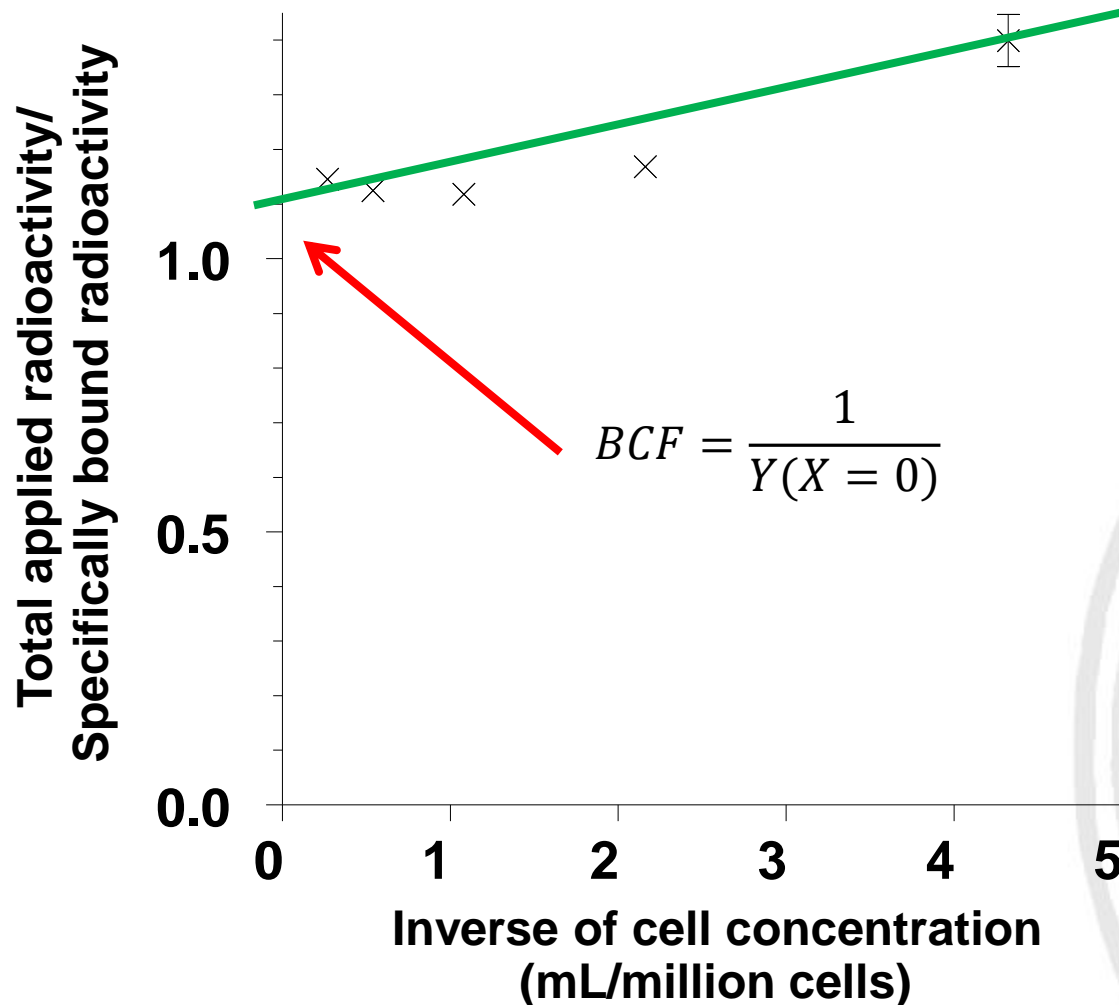
Fig. 1. Binding assay for the determination of the immunoreactive fraction of an ^{111}In -labeled T101 monoclonal antibody. The assay was set up using 6 concentrations of live NCI-H516 cells, in 1:2 dilutions from 3.3 to 0.1 million cells/ml, and the final concentration of ^{111}In -labeled T101 was 13 ng/ml. A shows a conventional plot of specific binding over total applied radioactivity, as a function of increasing cell concentration. B is a double inverse plot according to eq. (3) of the same data as in A, i.e., total applied radioactivity over specific binding, as a function of the inverse cell concentration. By means of linear extrapolation to the ordinate, the immunoreactive fraction can be determined for conditions representing infinite antigen excess.





Binding competent fraction

Increasing concentration of targeted **molecule**





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In vitro characterisation

Specificity:

Binding properties:

Cellular processing:

Binding

Internalization

Retention

Therapy effect





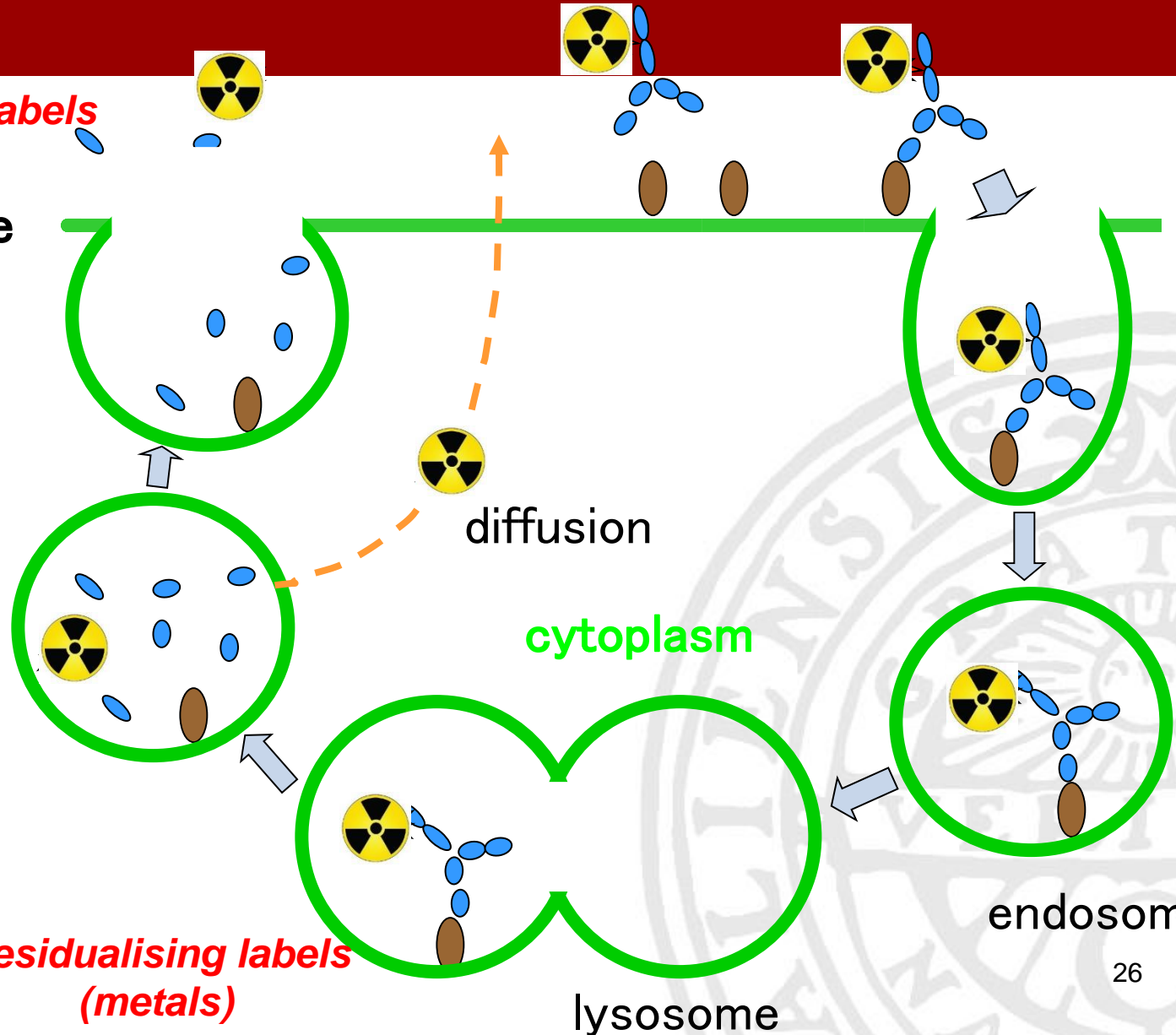
Internalization of radiolabeled proteins

externalization

internalization

*Non-residualising labels
(halogens)*

cell membrane



*Residualising labels
(metals)*

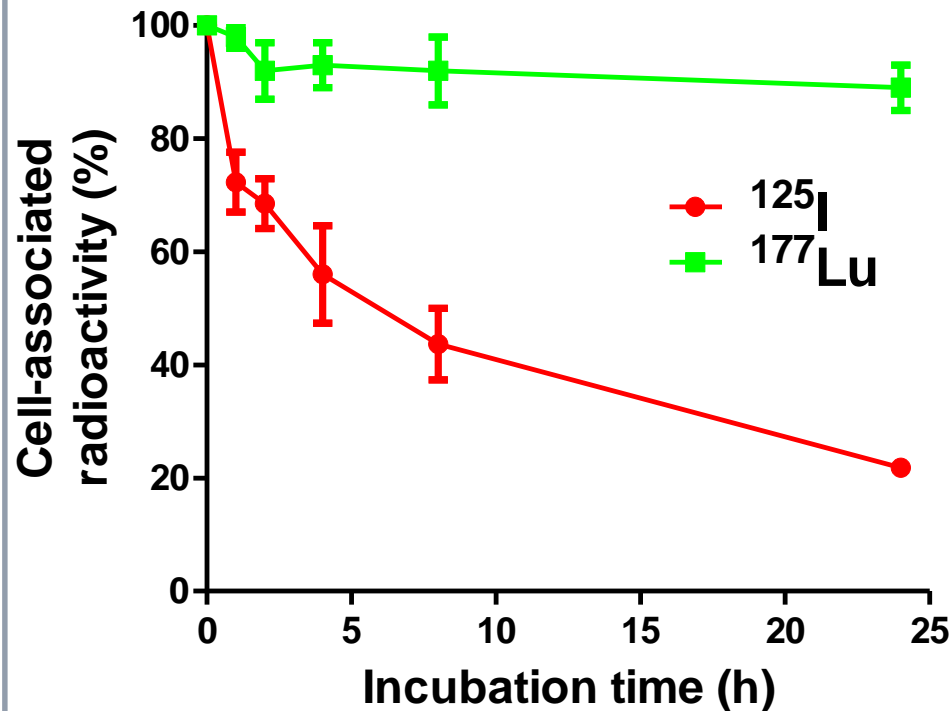
endosome

lysosome



Cellular retention of radionuclides

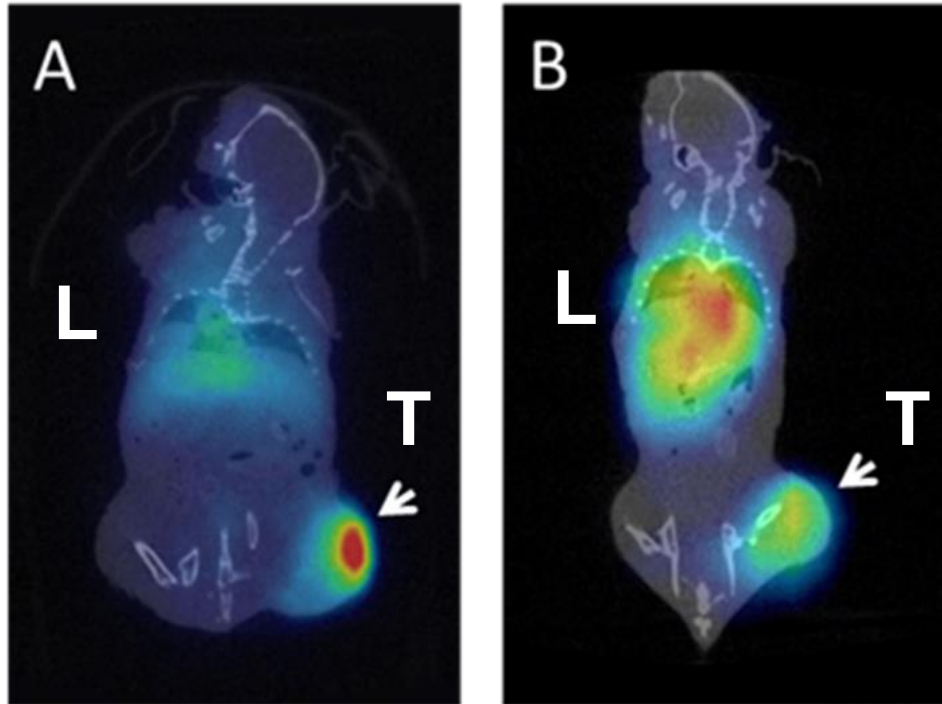
Processing of trastuzumab by breast cancer cells in vitro



The use of residualizing labels (metals) improves cellular retention of radionuclides delivered by antibodies because antibodies are internalized rapidly!



Retention in excretory organs



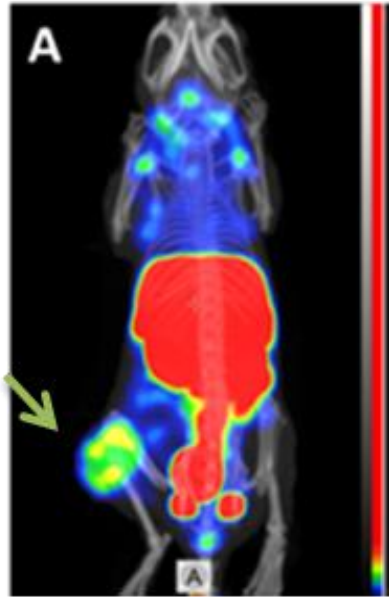
^{124}I : non-residualizing ^{111}In : residualizing

**Internalization in excretory organs is much faster
than in tumors**



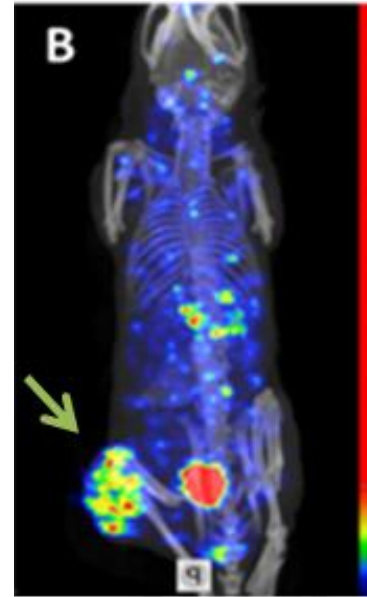
Same protein, different labels

**Residualizing
label ($^{99m}\text{Tc}(\text{CO})_3$)**



**Retention in tumor,
Retention in normal organs**

**Non-residualizing
label (^{125}I halogen)**



**Retention in tumor,
Wash-out from normal organs**

Scaffold proteins are *usually* internalized slowly by tumor cells



Comparison of labels

A **residualizing label** is trapped inside the cell after internalization and proteolysis of a labeled protein. It shows the location of protein uptake and catabolism that occurred **prior to the studied time point**



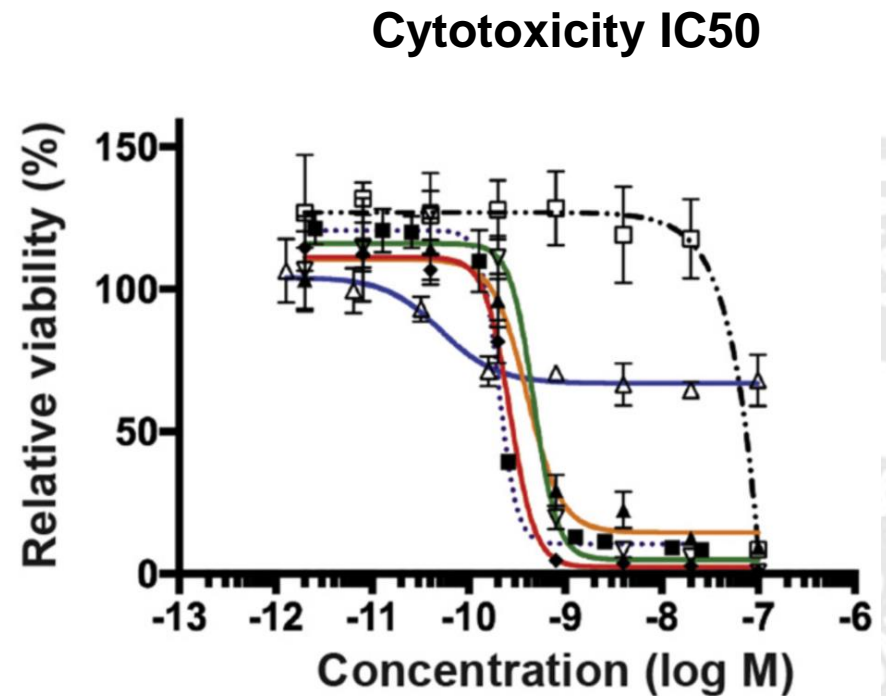
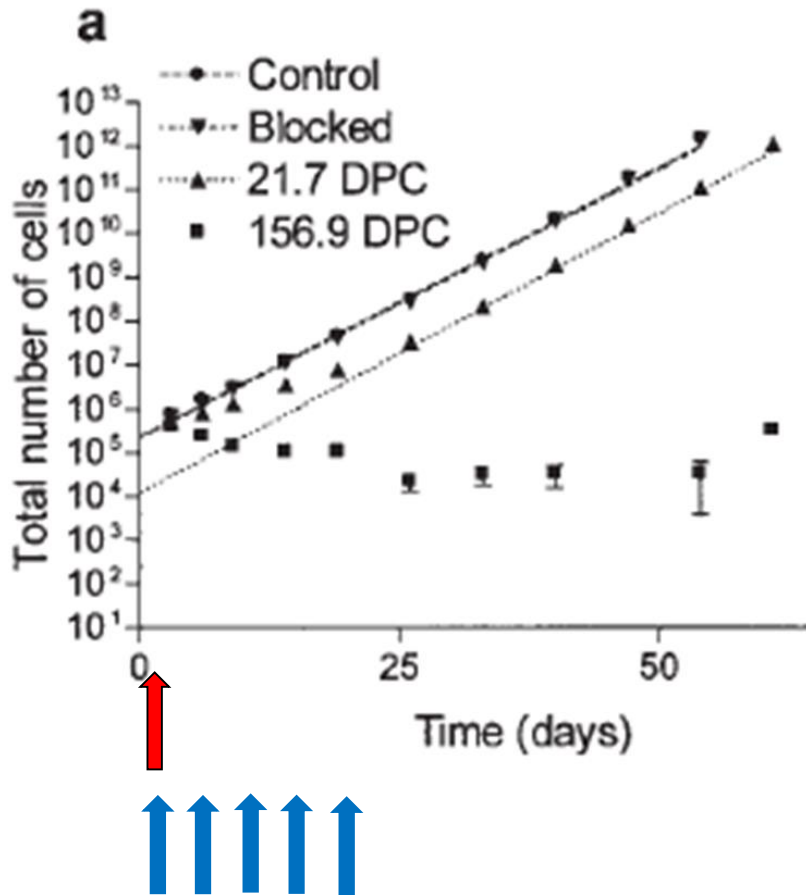
A **non-residualizing label**, i.e. its radiocatabolites diffuse from cells, shows the location of a labeled protein in the extracellular space/membrane **at the studied time point**



Comparison of these two types of label **provides information about protein localization** (intracellular space or cell membrane) and speed of internalization *in vivo*



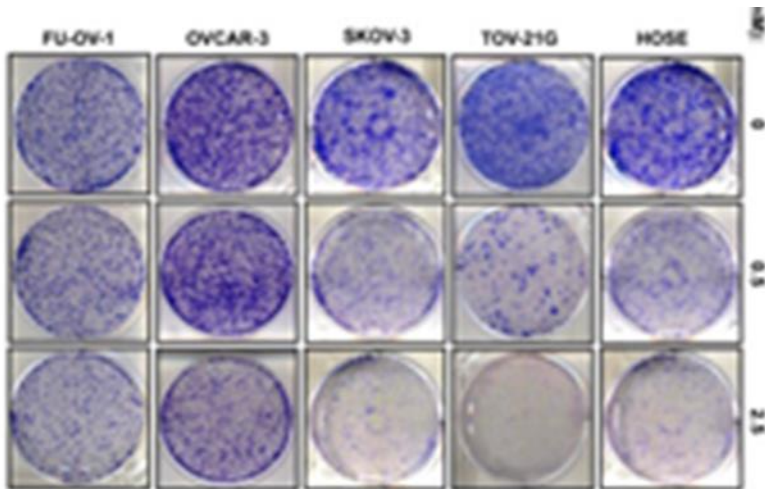
Therapy effect: growth curves



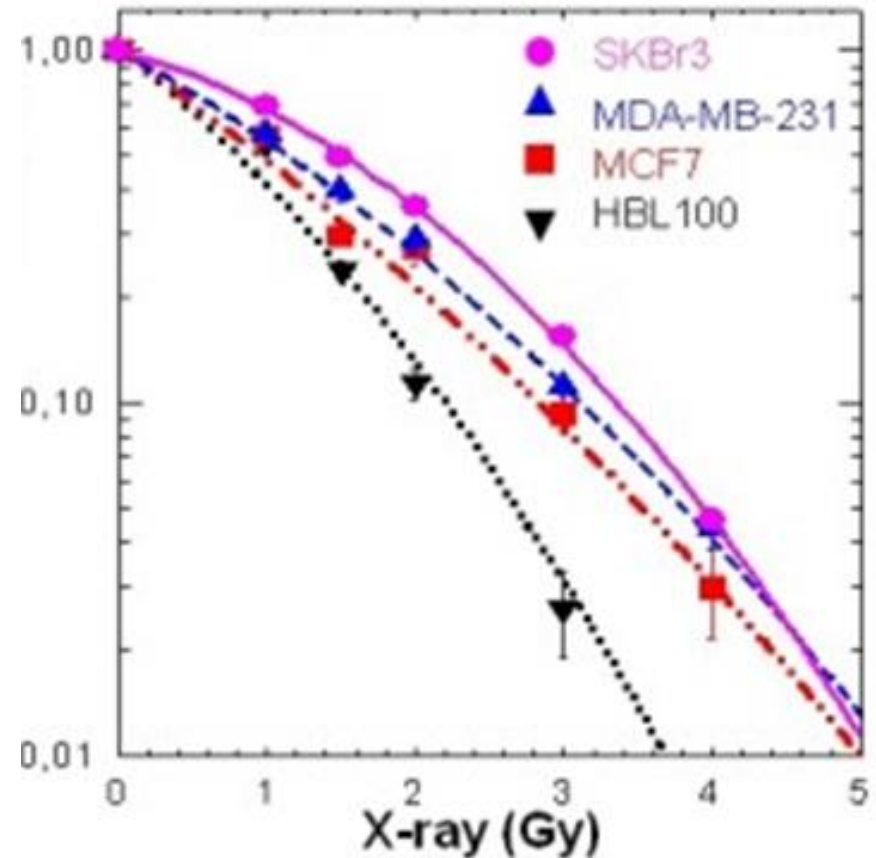


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Therapy effect: clonogenic survival assay



External beam radiation therapy





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In vivo characterisation

Normal biodistribution

Target specificity

Biodistribution in tumor bearing mice

Imaging (confirmation)

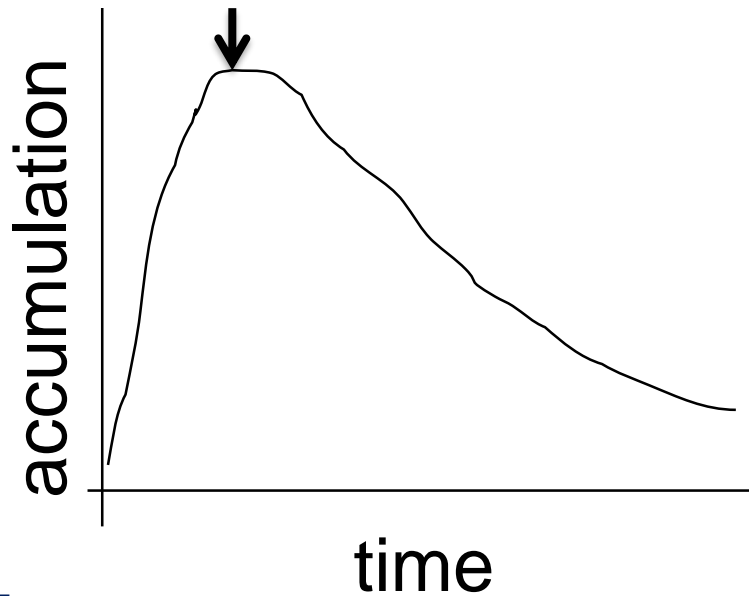
Dosimetry

Therapy

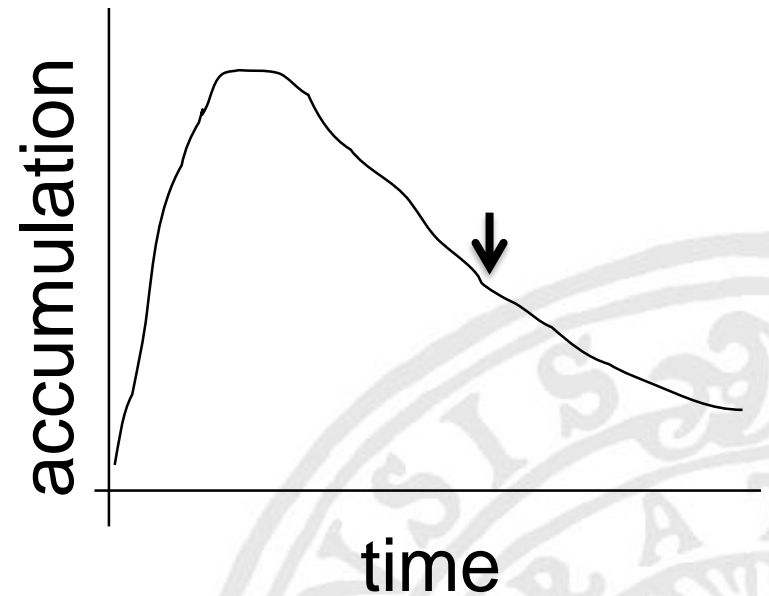




Biodistribution: time points



$T_{1/2}$ —
Observation



$T_{1/2}$ —
—

Case 1. Absolute accumulation

Example: small protein, 1 - 4 h

Case 2. Retention and wash out

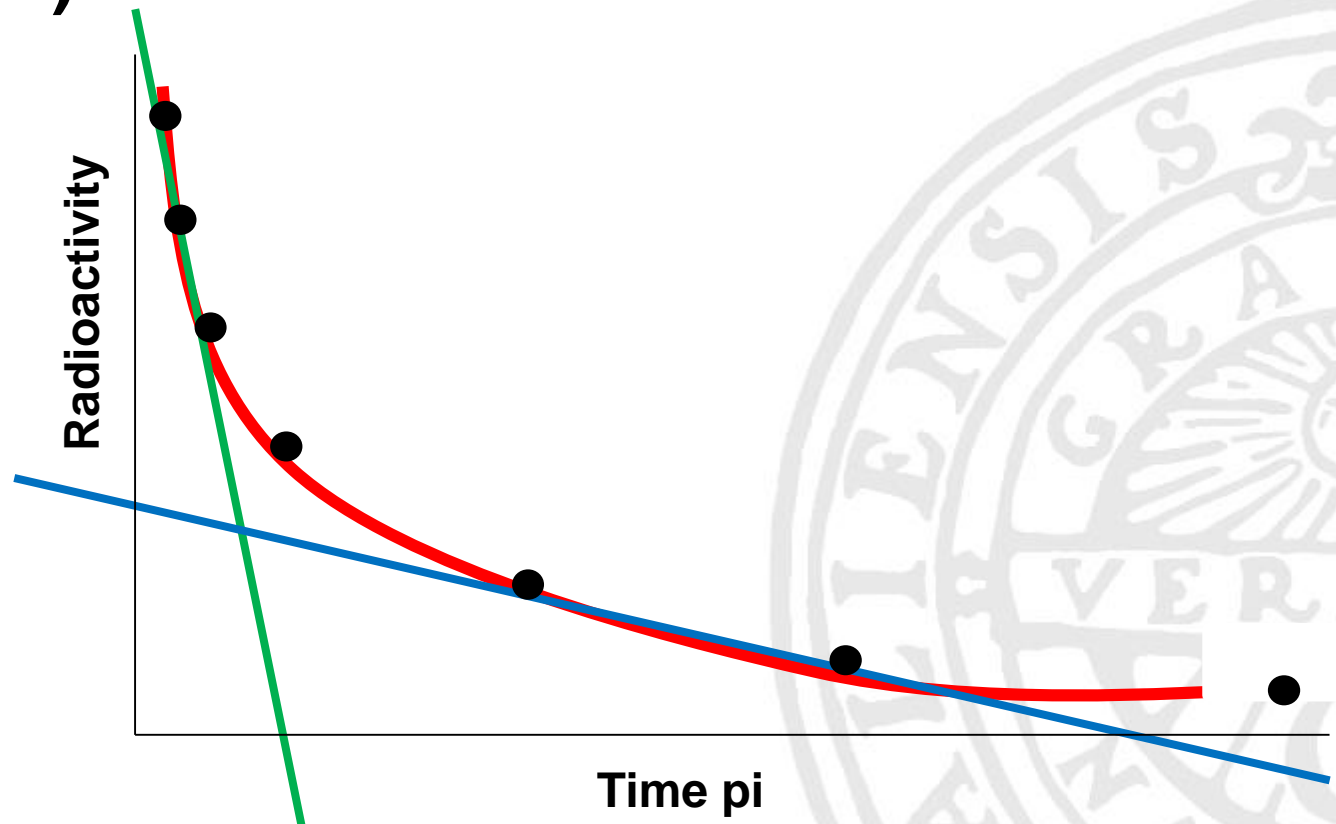
Example: small protein, 24 - 72 h



Normal biodistribution

Animals (without tumors)

Several time points (distribution phase + elimination)





Normal biodistribution

Animals (without tumors)

Several time points (distribution phase + elimination)

Organs should be studied

blood

excretory organs (liver, kidneys)

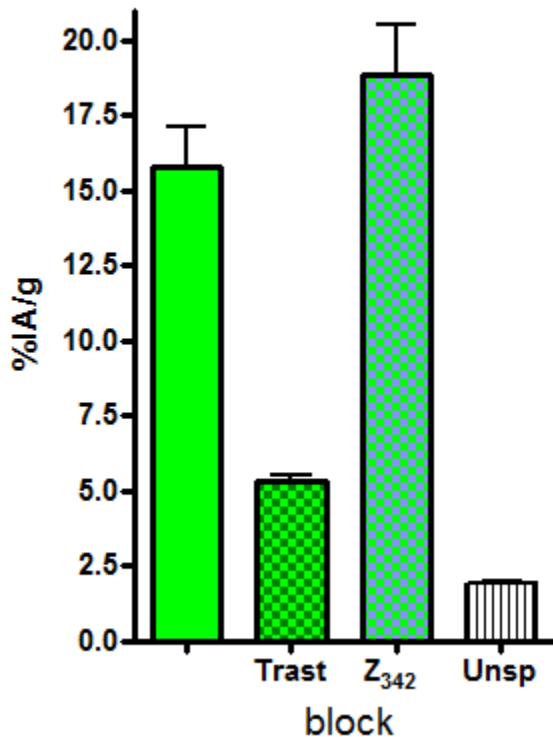
organs with normal target expression

organ with high accumulation of radiocatabolites

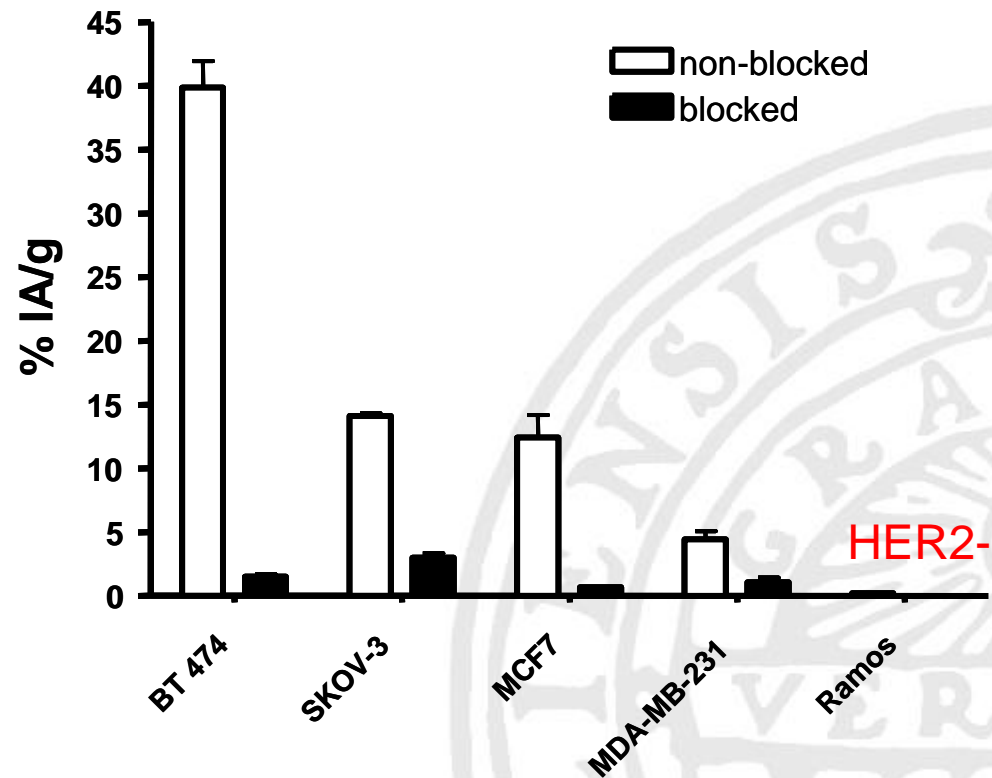


In vivo target specificity

Trastuzumab



HER2 targeting



1. Block by itself
2. Use a compound of same size but no specific binding

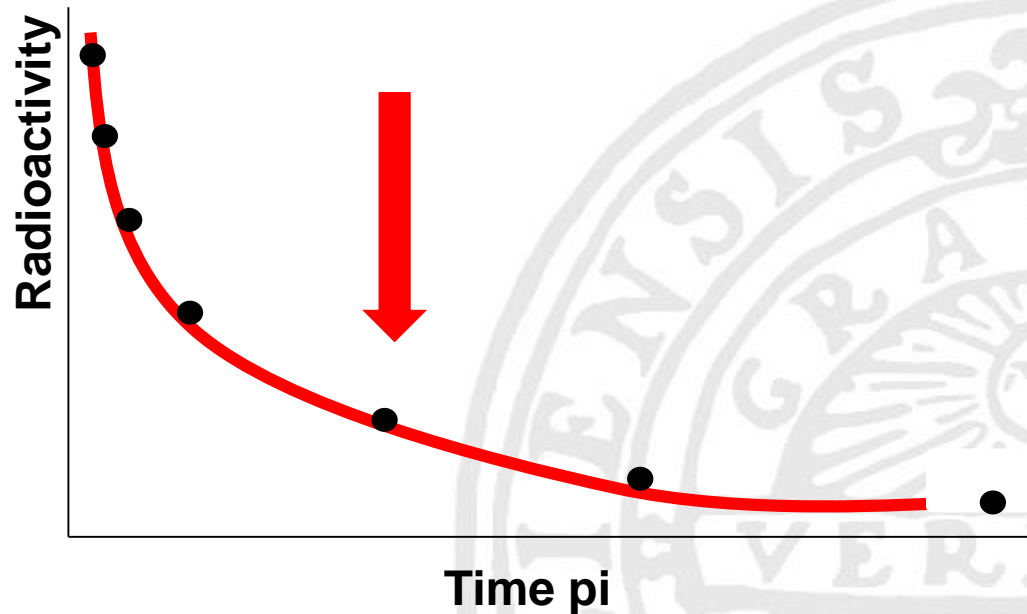
3. Several HER2+ or HER2- tumor models



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Biodistribution in tumor bearing mice

Could be one time point experiment (dedicated from normal biodistribution experiment)

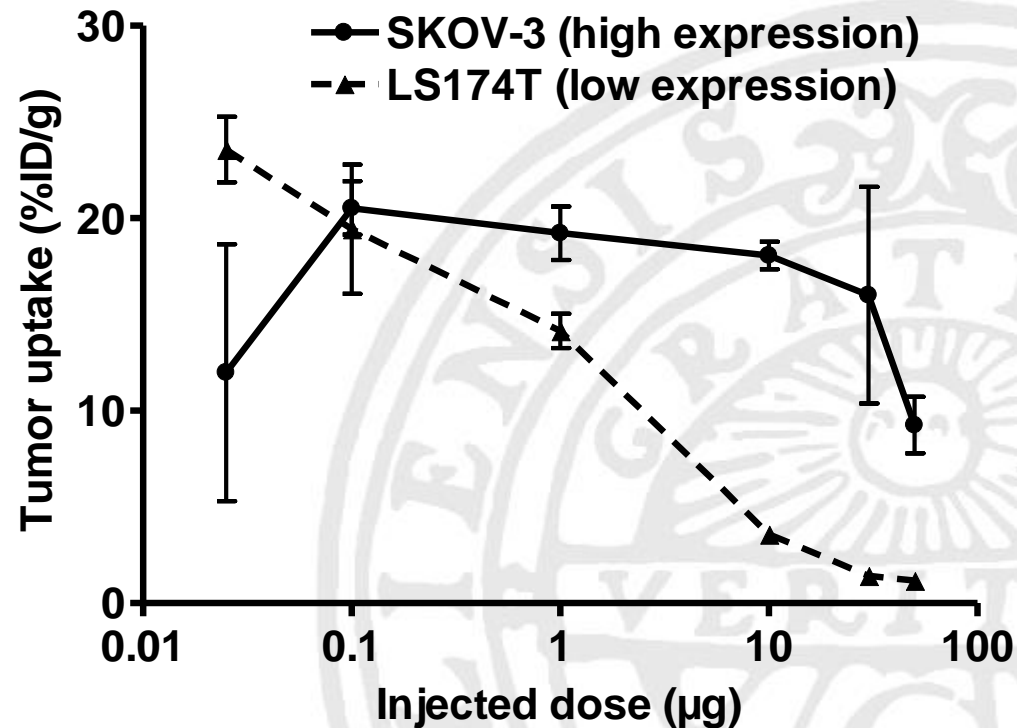
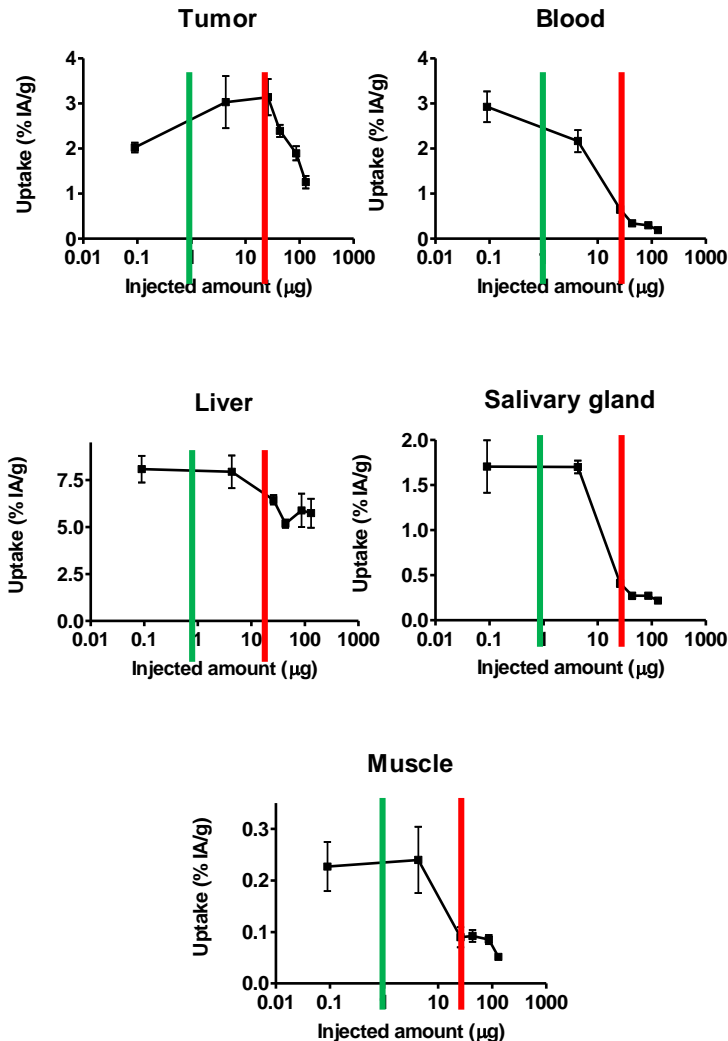




Biodistribution in tumor bearing mice

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Optimal injected specific radioactivity





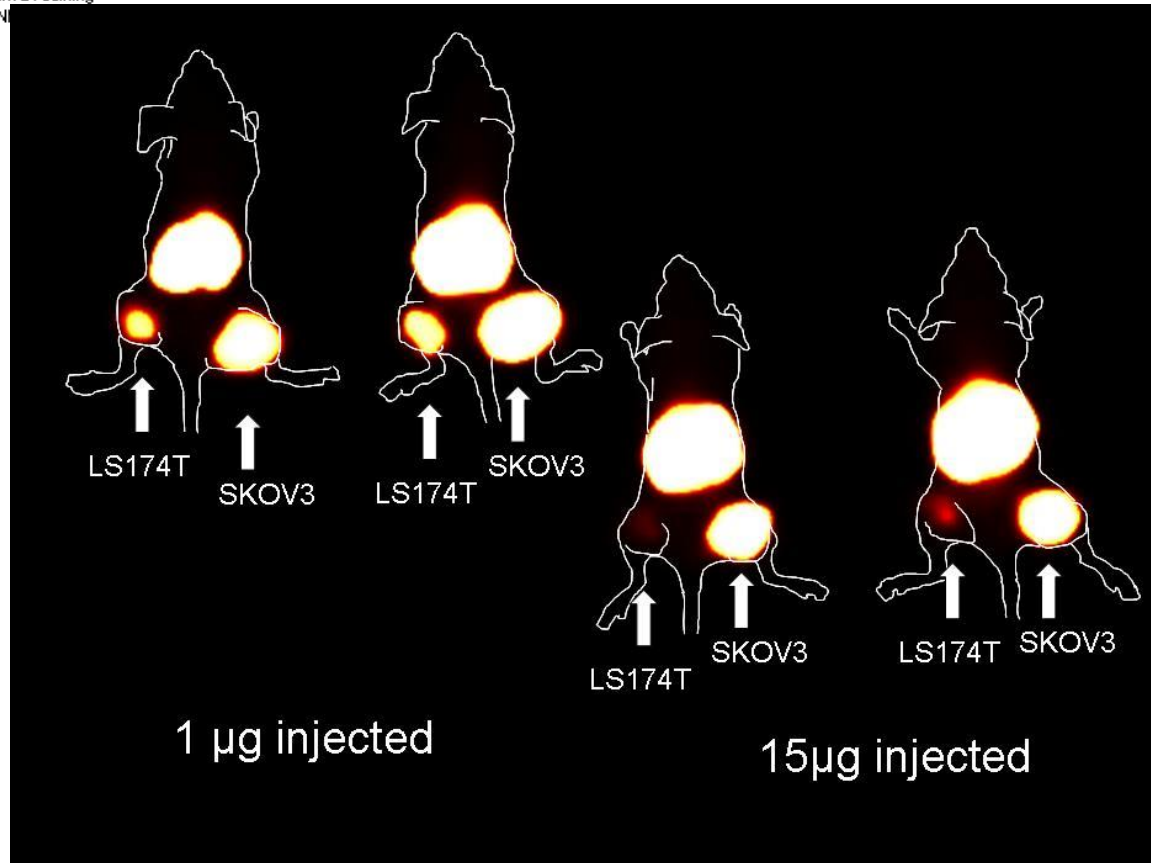
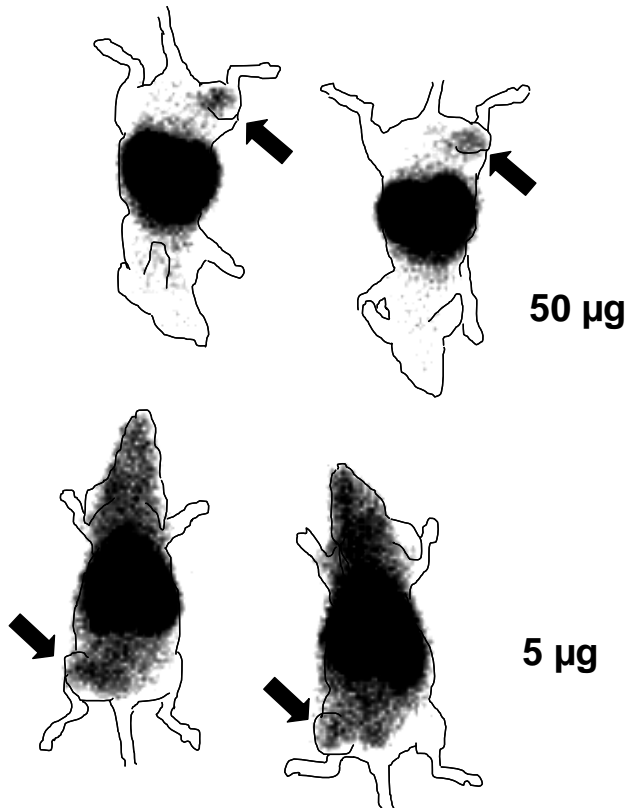
Biodistribution in tumor bearing mice

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Optimal injected specific radioactivity

1In_Z2377_DOSE_0
G_7

Akademiska Sjukhuset Uppsala, NM avdelning
N





Dosimetry

Imaging targeting molecule

needed if clinical trial is planned

multiple time points full scale

biodistribution in normal mice

Nuclide-based therapeutic targeting molecule

needed for planning of therapy experiment

multiple time point full scale biodistribution in
tumor bearing mice

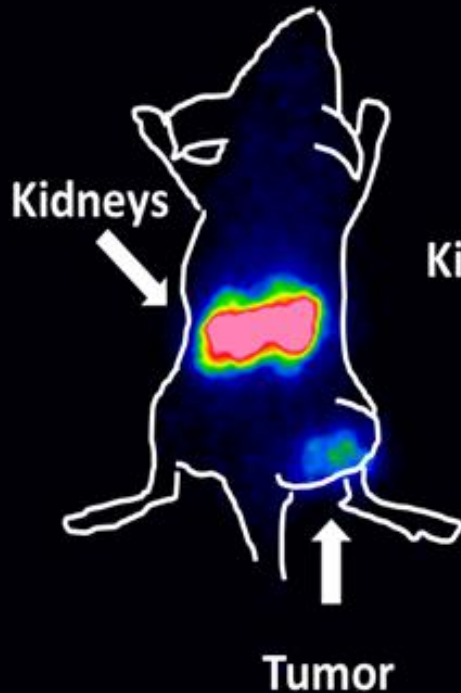
Area under curve (AUC) for tumors >> for
healthy organs



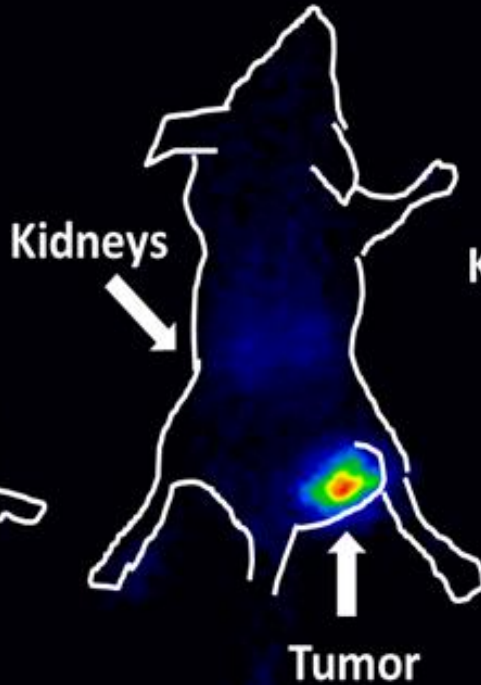
Affibody-Based PNA Pretargeting

(1 h p.i.)

$^{111}\text{In-DOTA-Z}_{\text{HER2:K58}}$
Direct labeling



$Z_{\text{HER2:342-HP1+}}$
 $^{111}\text{In-DOTA-HP2}$
Pretargeting



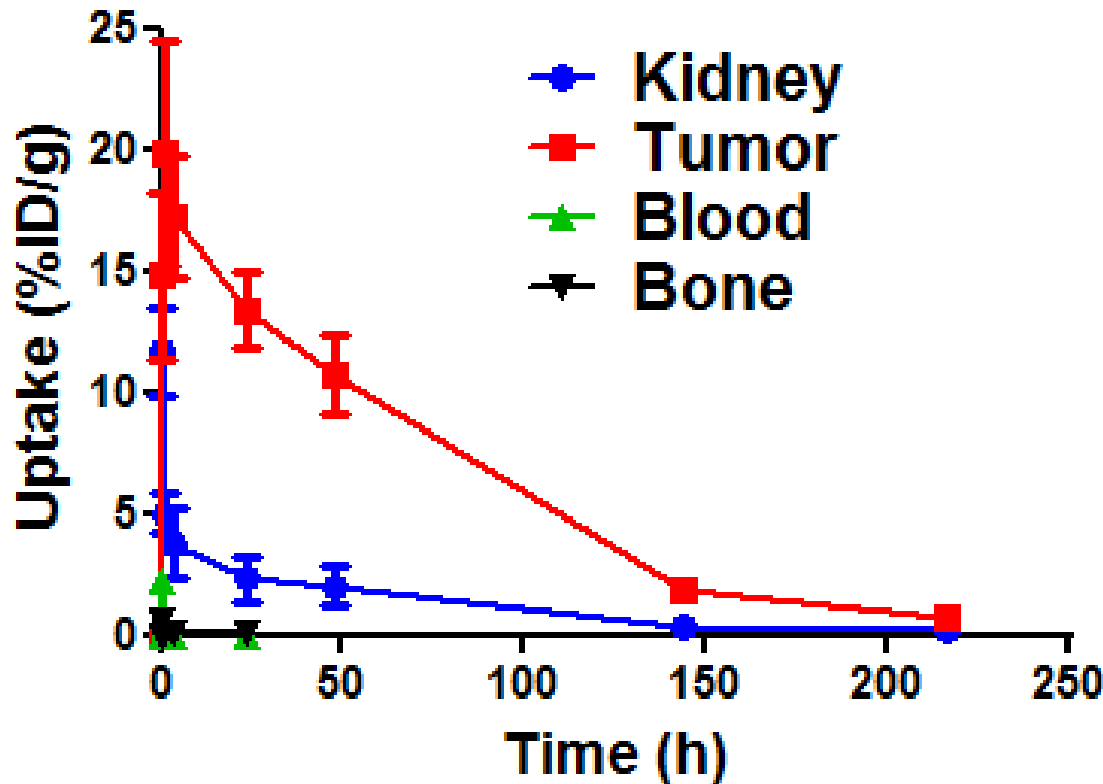
$^{111}\text{In-DOTA-HP2}$
Negative control





Dosimetry

Residence of ^{177}Lu (decay uncorrected)



AUC	ratio
268	5
1362	1
1.9	716
1.2	1048

Calculation of Area Under Curve for Dosimetry Evaluation



Experimental therapy

Should be carefully planned (expensive, time consuming)

appropriate control groups

vehicle (PBS, saline)

unspecific molecule with similar kinetics

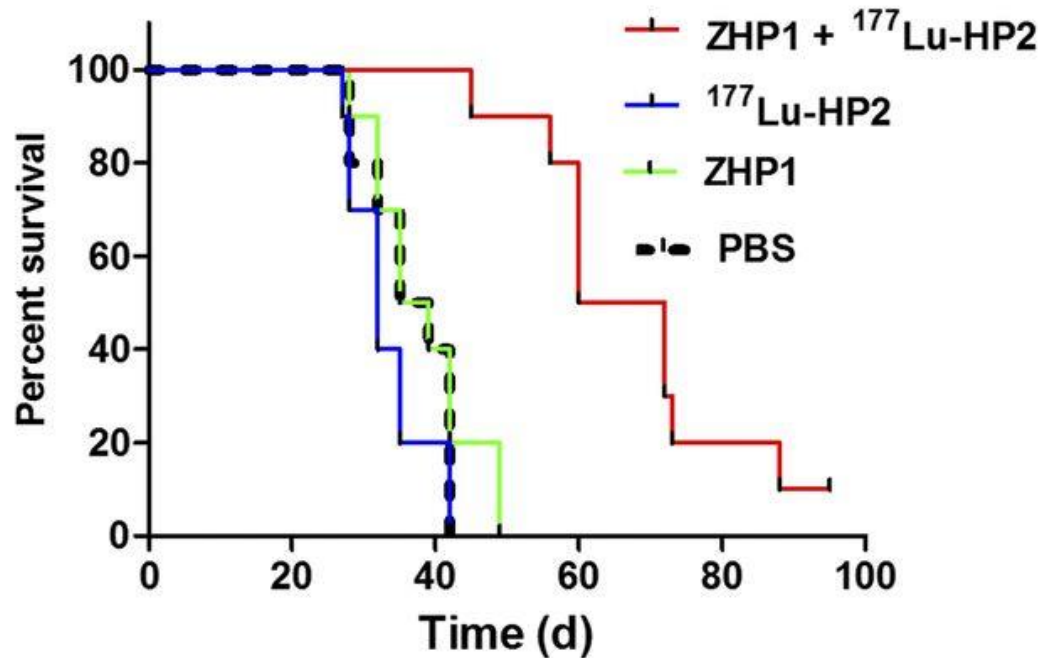
un-labeled targeting molecule

number of animals per group

end point



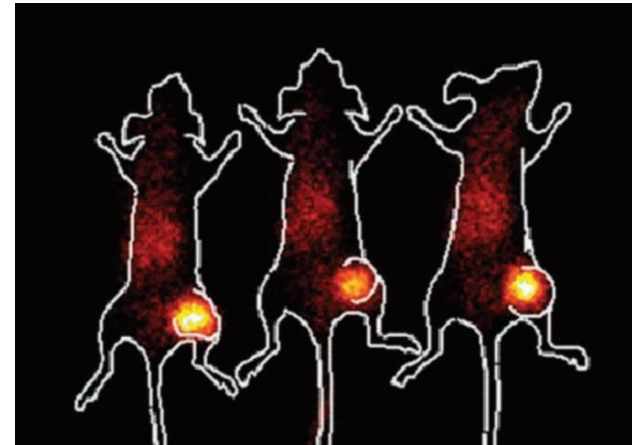
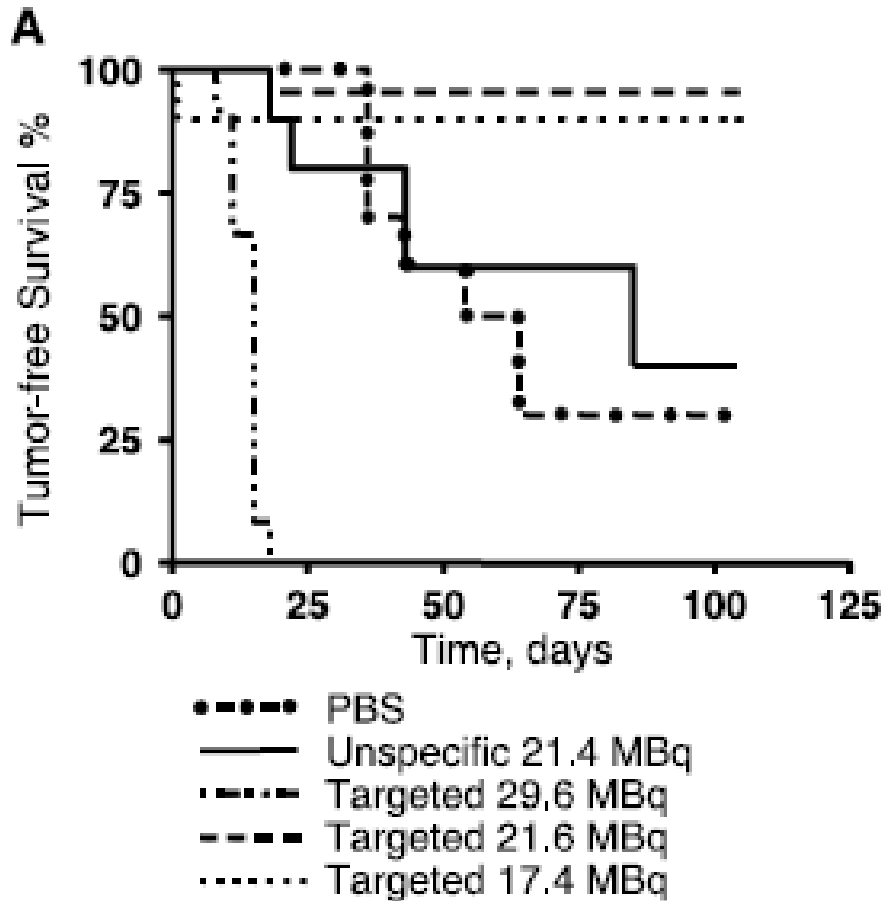
Radionuclide Therapy of HER2+ Xenografts Using Affibody-Based PNA Pretargeting



6 cycles of radionuclide therapy with $^{177}\text{Lu-HP2}$ doubled median survival of mice (66 d. vs 37 d.)



Radionuclide therapy with ^{177}Lu -affibody-ABD prevented formation of tumors in mice



Tolmachev et al. *Cancer Res.* 2007; 67(6):2773–82.

Orlova et al. *J. Nucl. Med.* 2013; 54: 961–968



Pre-clinical evaluation of radiolabeled proteins

