



Characterizing In vitro Synergy using the Ray design methodology

Application to an Oncology combination study

Sophie Ruquet, Noëlle Boussac
SANOFI-AVENTIS
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sanofi aventis

L'essentiel c'est la santé.



Outline

- Synergy definition and Loewe additivity model
- Ray design methodology applied to an Oncology combination study
 - Context and preliminary data
 - Experimental ray design
 - Statistical analysis and modeling
 - Model fitting and results validation
 - Conclusion
- Ray design methodology: pros and cons



Synergy – definition (1/2)

- **Definition <pharmacology, physiology>**: The interaction of two or more treatments such that their combined effect is greater than the sum of the individual effects observed when each treatment is administered alone
- **Loewe additivity model (Loewe and Muischnek, 1926)**
 - Most suitable reference model
 - Reasoning at fixed effect: in a synergistic mixture, lower concentrations of the two products are needed to obtain a given effect, in comparison with additive situation
- **Equation for Loewe additivity model between products A and B**

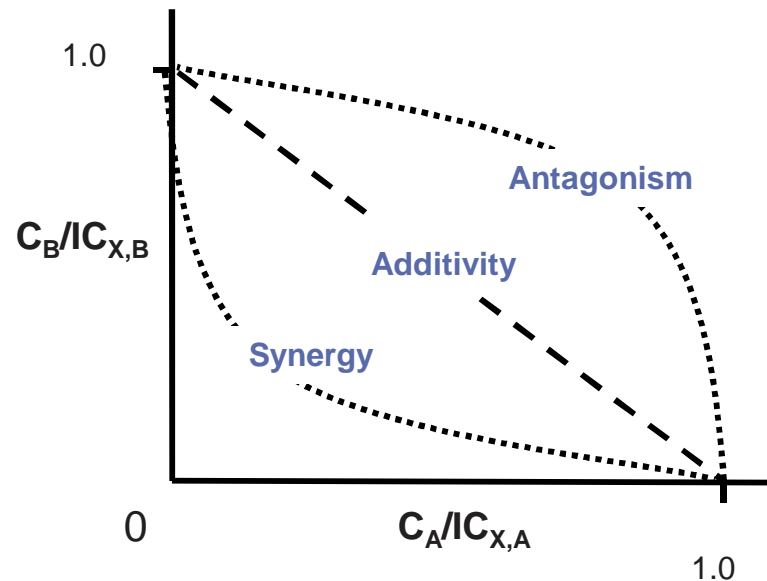
$$\frac{C_A}{IC_{X,A}} + \frac{C_B}{IC_{X,B}} = 1$$

- C_A, C_B : concentrations of each product in the mixture necessary to obtain X% of effect
- $IC_{X,A}, IC_{X,B}$: concentrations of products A and B necessary to obtain X% of effect for each product alone (often relative IC50, concentration to obtain 50% of the delta between the maximum and the minimum effects)



Synergy – definition (2/2)

- The left-hand term $K_i = \frac{C_A}{IC_{X,A}} + \frac{C_B}{IC_{X,B}}$ of the equation is significantly:
- Inferior to 1 in case of synergy
 - Superior to 1 in case of antagonism

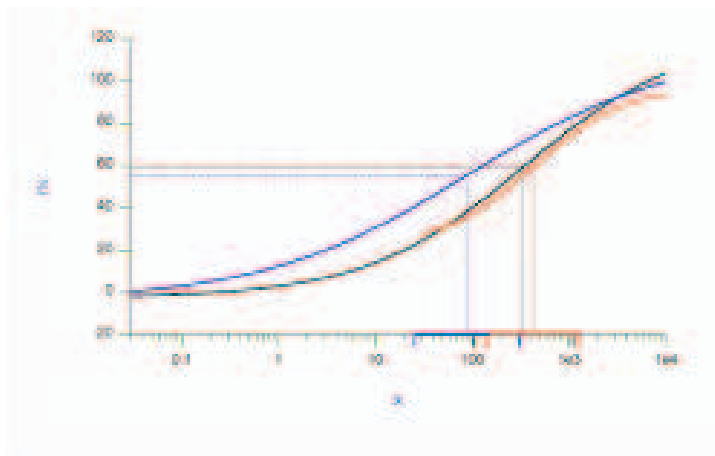




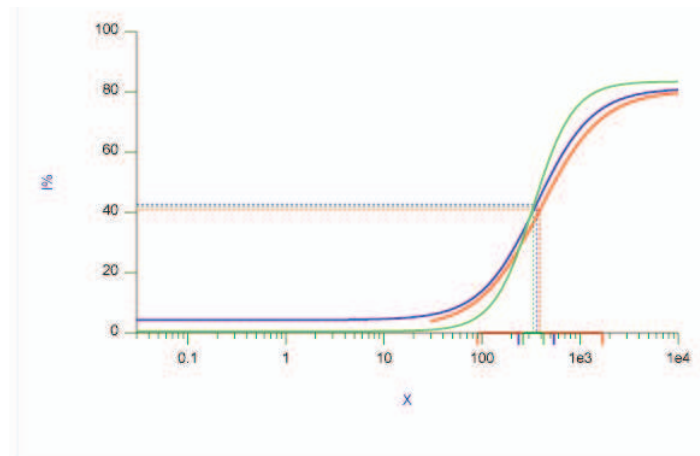
Study context and preliminary data

- Objective: study the In vitro combination of two anticancer agents, Prod.A and Prod.B, to detect their possible synergy on a given cancer cell line
- Parameter of interest: Percentage of inhibition of cancer cells growth
- Preliminary results on each product alone: relative IC50s, minimum and maximum concentrations

Product A: IC50=300nM
Min=0.03nM
Max=10000nM



Product B: IC50=300nM
Min=3nM
Max=10000nM





Experimental Ray design (1/4)

- Each ray design contains at least 5 rays:
 - Two rays corresponding to each product alone
 - Other rays i consisting of couples of concentrations of products A and B, in a given proportion $c_i = C_{\text{Prod.B}}/C_{\text{Prod.A}}$, constant for each ray
 - Each couple of concentrations in duplicates, at least 6-7 successive dilutions by ray within the minimum and maximum concentrations range of each product
- The synergy zone is covered in a symmetric way from the equipotent ray, where $c_i = IC50_{\text{Prod.B}}/IC50_{\text{Prod.A}}$ and products are equally represented considering their respective potency
- At least three independent experiments (3 Ray designs) performed to ensure robustness of results

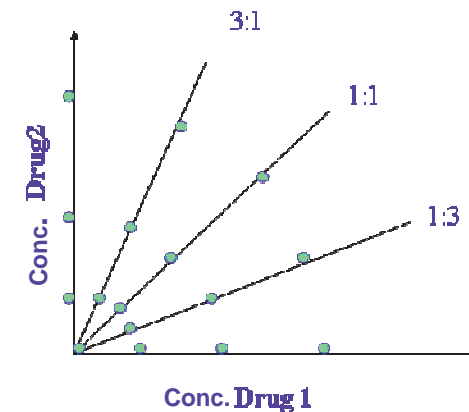


Experimental Ray design (2/4)

- For each ray, each proportion $c_i = C_{\text{Prod.B}}/C_{\text{Prod.A}}$ translated into unit of effect of each product alone considering their respective $IC50s'$ values, using the effective fraction

$$f: f_i = \frac{1}{c_i \rho + 1} \quad \text{where} \quad \rho = \frac{IC50_{\text{Prod.A}}}{IC50_{\text{Prod.B}}} \quad \text{is the relative potency of the two products}$$

- Ex1: $f=0.5$, effective equipotent ray ($f \in]0,1[$)
- Ex2: $f=0.75$, ray where Prod.A is 3 times more represented than Prod.B considering their relative potency, also called ray « 3 for 1 »



- In the study, $\rho = \frac{300\text{nM}}{300\text{nM}} = 1$
- With 5 rays, taking the two products alone and 3 rays with $f=0.75$ (ray 3 for 1), $f=0.5$ (equipotent ray) and $f=0.25$ (ray 1 for 3) permits to cover equally all the synergy zone



Experimental Ray design (3/4)

Example: Ray design for Experiment 1 (concentrations in nM)

Ray 1 : Product A alone

| | | | | | | | | | | |
|---------|------|-----|-----|----|----|---|---|-----|-----|------|
| Mixture | 1000 | 300 | 100 | 30 | 10 | 3 | 1 | 0.3 | 0.1 | 0.03 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 | 3 | 1 | 0.3 | 0.1 | 0.03 |
| Prod.B | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |

Ray 2 (1 for 30, $f=0.03$) :
300nM (Prod.A) + 10000nM (Prod.B)

| | | | | | | | | | |
|---------|-------|-------|------|------|-----|-----|----|------|-----|
| Mixture | 21000 | 10300 | 3100 | 1030 | 310 | 103 | 31 | 10.3 | 3.1 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 | 3 | 1 | 0.3 | 0.1 |
| Prod.B | 20000 | 10000 | 3000 | 1000 | 300 | 100 | 30 | 10 | 3 |

Ray 3 (1 for 10, $f=0.09$) :
300nM (Prod.A) + 3000nM (Prod.B)

| | | | | | | | | |
|---------|-------|------|------|-----|-----|----|----|-----|
| Mixture | 10100 | 3300 | 1100 | 330 | 110 | 33 | 11 | 3.3 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 | 3 | 1 | 0.3 |
| Prod.B | 10000 | 3000 | 1000 | 300 | 100 | 30 | 10 | 3 |



Experimental Ray design (4/4)

Ray 4 (1 for 3, $f=0.25$) :
300nM (Prod.A) + 1000nM (Prod.B)

| | | | | | | | |
|---------|------|------|-----|-----|-----|----|---|
| Mixture | 4000 | 1300 | 400 | 130 | 400 | 13 | 4 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 | 3 | 1 |
| Prod.B | 3000 | 1000 | 300 | 100 | 30 | 10 | 3 |

Ray 5 (1 for 1, $f=0.50$) :
300nM (Prod.A) + 300nM (Prod.B)

| | | | | | | |
|---------|------|-----|-----|----|----|---|
| Mixture | 2000 | 600 | 200 | 60 | 20 | 6 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 | 3 |
| Prod.B | 1000 | 300 | 100 | 30 | 10 | 3 |

Ray 6 (3 for 1, $f=0.75$) :
1000nM (Prod.A) + 300nM (Prod.B)

| | | | | | |
|---------|------|-----|-----|----|----|
| Mixture | 1300 | 400 | 130 | 40 | 13 |
| Prod.A | 1000 | 300 | 100 | 30 | 10 |
| Prod.B | 300 | 100 | 30 | 10 | 3 |

Ray 7 : Product B alone

| | | | | | | | | | |
|---------|-------|-------|------|------|-----|-----|----|----|---|
| Mixture | 20000 | 10000 | 3000 | 1000 | 300 | 100 | 30 | 10 | 3 |
| Prod.A | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Prod.B | 20000 | 10000 | 3000 | 1000 | 300 | 100 | 30 | 10 | 3 |



Statistical analysis of the Ray design

- For each experiment, global modelling of all rays together, each ray following a 4-parameter concentration-effect (inhibition) logistic curve

- For each experiment and from this global model:
 - Estimation of the experimental values of the IC50s of Prod.A and Prod.B alone

 - Estimation of the experimental values of « f »

 - Estimation of the Loewe additivity index K_i for each ray i ($i=2..6$)



Modeling the Ray design

Concentration-effect model

- A 4-parameter logistic model for each ray:

$$E[Y] = E_{\min} + \frac{E_{\max} - E_{\min}}{1 + \exp(-m \times \ln(\frac{\text{Conc}}{\text{IC50}}))}$$

Maximum inhibition effect $\rightarrow E_{\max}$
 Minimum inhibition effect $\rightarrow E_{\min}$
 Total concentration : $C_A + C_B$
 Expected value of the percentage of inhibition of the product alone or the mixture $\rightarrow E[Y]$
 Slope of the conc.-effect curve $\rightarrow m$
 IC50 of the ray $\rightarrow \text{IC50}$

- Generalization to all rays i , $i=1, \dots, 5$ with different parameters E_{\max} , E_{\min} , m , IC50 for each ray:

Global model:

$$E[Y] = E_{\min_i} + \frac{E_{\max_i} - E_{\min_i}}{1 + \exp(-m_i \ln(\frac{\text{Conc}_i}{\text{IC50}_i}))}$$



Measurement of interaction between drugs (1/2)

- K_i value, $i=2, 3, 4$, permits to measure the relationship (additivity, antagonism or synergy) between the 2 tested products A and B for Ray i:

$$K_i = \frac{IC50_i (IC50_A + c_i IC50_B)}{IC50_A IC50_B (1 + c_i)}$$

where

- $c_i = C_B / C_A$ for Ray i
- $IC50_A$, IC50 for product A alone (ray 1)
- $IC50_B$, IC50 for product B alone (ray 5)
- $IC50_i$, with for rays $i=2, i=3$ and $i=4$, $c_i = C_B / C_A$

} Obtained by the
global model



Measurement of interaction between drugs (2/2)

- K_i value is obtained with its corresponding CI95%
- K_i value is significantly:
 - Equal to 1 in case of additivity (CI95% contains '1')
 - Inferior to 1 in case of synergy (CI95% is strictly inferior to '1')
 - Superior to 1 in case of antagonism (CI95% is strictly superior to '1')
- E_{min_i} , E_{max_i} , m_i , $IC50_i$ and K_i estimates are obtained with NLMixed procedure



Fitting the global concentration-effect model

- **NLMixed SAS software procedure**
- **Parameters initialization: use of estimations of parameters obtained from the fitting of each curve separately**
- **For each experiment, simultaneous estimation of adjusted curves parameters: E_{min_i} , E_{max_i} , $IC50_i$, slope m_i and K_i with $i=1, \dots, 5$**
- **Selection of the best model**
 - *First step*: full model with E_{min_i} , E_{max_i} and m_i specific to each ray
 - *Second step*: for this model, test of equality of the parameters E_{min_i} , E_{max_i} and m_i for all rays
 - *Third step*: new model in which the previously significant parameter(s) is(are) considered common to all rays



Results validation

| EXP. 2 | Preliminary values | Experimental values |
|-------------|--------------------|---------------------|
| IC50 Prod.A | 300 | 200.23 |
| IC50 Prod.B | 300 | 165.80 |
| f Ray 2 | 0.23 | 0.20 |
| f Ray 3 | 0.50 | 0.45 |
| f Ray 4 | 0.77 | 0.73 |
| f Ray 5 | 0.91 | 0.89 |
| f Ray 6 | 0.97 | 0.96 |

Conclusion domain similar to the expected one

| EXP. 1 | Preliminary values | Experimental values |
|-------------|--------------------|---------------------|
| IC50 Prod.A | 300 | 106.78 |
| IC50 Prod.B | 300 | 217.73 |
| f Ray 2 | 0.03 | 0.06 |
| f Ray 3 | 0.09 | 0.17 |
| f Ray 4 | 0.25 | 0.40 |
| f Ray 5 | 0.50 | 0.67 |
| f Ray 6 | 0.75 | 0.86 |

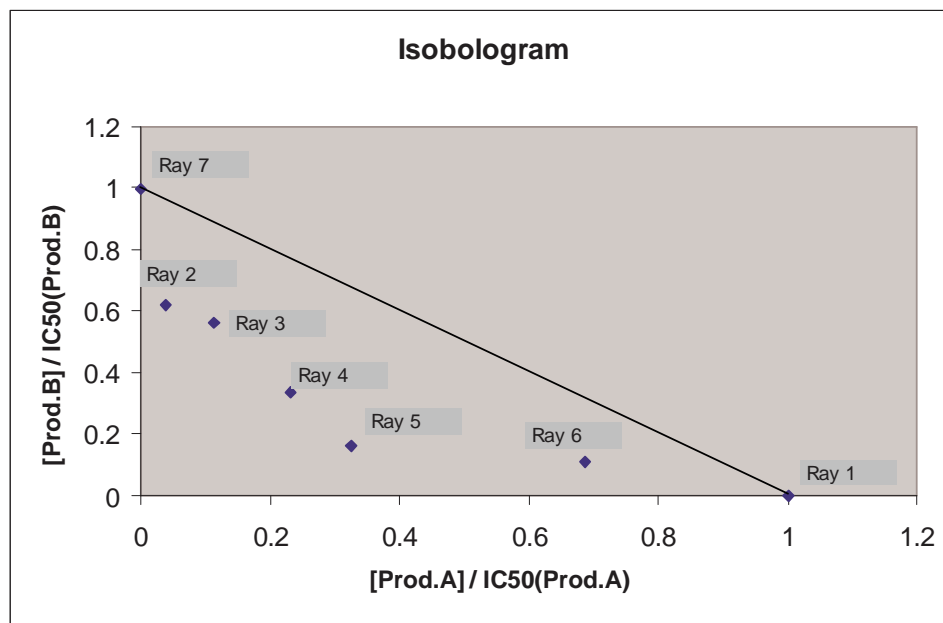
Right shift of the conclusion domain, f values higher than expected

| EXP. 3 | Preliminary values | Experimental values |
|-------------|--------------------|---------------------|
| IC50 Prod.A | 300 | 87.74 |
| IC50 Prod.B | 300 | 729.82 |
| f Ray 2 | 0.23 | 0.71 |
| f Ray 3 | 0.50 | 0.89 |
| f Ray 4 | 0.77 | 0.97 |

Shift to the domain where Prod.A is 3 times more represented than Prod.B considering their relative potency



Statistical results: Experiment 1



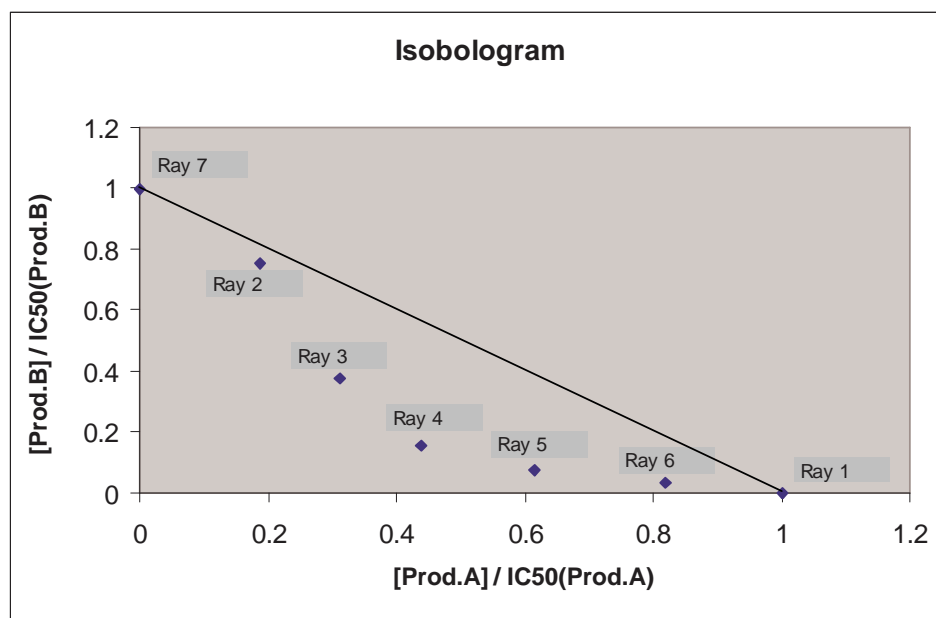
| EXP. 1 | Experimental f values | Ki values and 95%CI | Conclusion |
|--------|-----------------------|------------------------------|------------|
| Ray 2 | 0.06 | 0.6585 [0.4641; 0.8528] (*) | Synergy |
| Ray 3 | 0.17 | 0.6756 [0.4802; 0.8709] (*) | Synergy |
| Ray 4 | 0.40 | 0.5702 [0.3656; 0.7748] (*) | Synergy |
| Ray 5 | 0.67 | 0.4865 [0.2316; 0.7414] (*) | Synergy |
| Ray 6 | 0.86 | 0.8006 [0.1445; 1.4568] (NS) | Additivity |

(*): K is significantly different from 1 at 0.05 level

NS: Non significant



Statistical results: Experiment 2



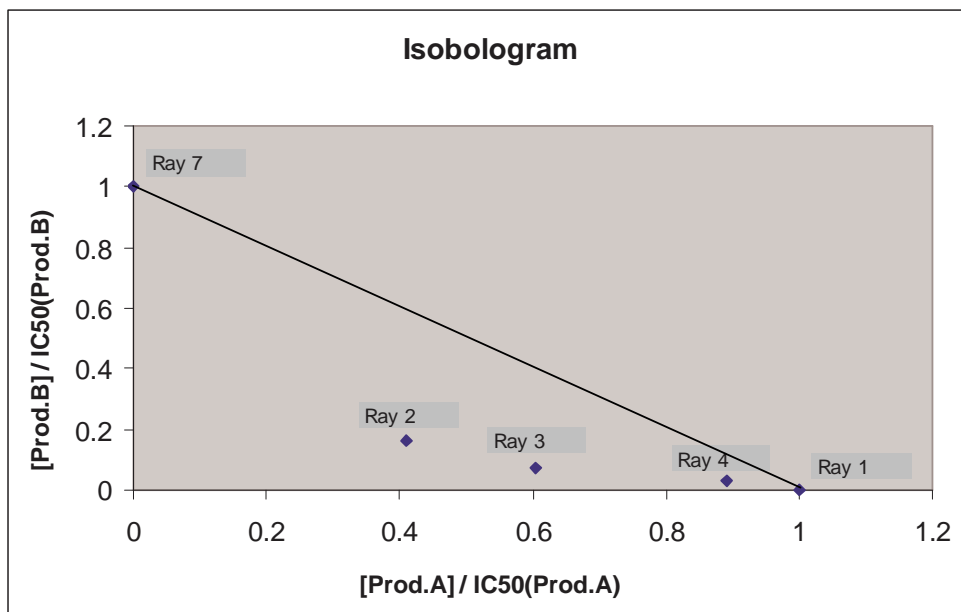
| EXP. 2 | Experimental f values | Ki values and 95%CI | Conclusion |
|--------|-----------------------|------------------------------|------------|
| Ray 2 | 0.20 | 0.9441 [0.7094; 1.1788] (NS) | Additivity |
| Ray 3 | 0.45 | 0.6906 [0.4794; 0.9018] (*) | Synergy |
| Ray 4 | 0.73 | 0.5987 [0.3731; 0.8243] (*) | Synergy |
| Ray 5 | 0.89 | 0.6891 [0.3615; 1.0166] (NS) | Additivity |
| Ray 6 | 0.96 | 0.8523 [0.3127; 1.3918] (NS) | Additivity |

(*): K is significantly different from 1 at 0.05 level

NS: Non significant



Statistical results: Experiment 3



| EXP. 3 | Experimental f values | Ki values and 95%CI | Conclusion |
|--------|-----------------------|------------------------------|------------|
| Ray 2 | 0.71 | 0.5746 [0.2503; 0.8989] (*) | Synergy |
| Ray 3 | 0.89 | 0.6770 [0.1806; 1.1734] (NS) | Additivity |
| Ray 4 | 0.97 | 0.9226 [0.2104; 1.6348] (NS) | Additivity |

(*): K is significantly different from 1 at 0.05 level

NS: Non significant



Global conclusion

- Summary of the experimental f values and associated conclusions : synergy (syn), additivity (add) or antagonism (ant)

| Experimental f values | | |
|-----------------------|-------------------|-------------------|
| EXP.1 | EXP.2 | EXP.3 |
| 0.06 (syn) | 0.20 (add) | 0.71 (syn) |
| 0.17 (syn) | 0.45 (syn) | 0.89 (add) |
| 0.40 (syn) | 0.73 (syn) | 0.97 (add) |
| 0.67 (syn) | 0.89 (add) | |
| 0.86 (add) | 0.96 (add) | |

- For these three experiments, synergy between Prod.A and Prod.B is observed in the three first quarters of the domain (except for Exp2, f=0.20)
- Additivity between Prod.A and Prod.B is observed for higher f values
- On the tested rays, 'synergistic' domain : relative concentrations of Prod.A and Prod.B so that Prod.A is never more than 3 times more represented than Prod.B considering their relative potency



Ray design modeling methodology - Pros & cons

- **Adapted to the biological mechanism of synergy by:**
 - taking into account the relative proportions of products
 - allowing to consider the specificity of each ray
- **Less trial- and time-consuming than full designs like grid designs**
- **Permit an accurate research of the synergy zone based on the proportion condition of the two compounds**

- **Can be only applied on in-vitro multiple-dose experiments with at least 6-7 different doses by ray**
- **Require a good knowledge of the studied products : accurate products IC50s, max and min effects estimations**
- **Adequacy between the expected and observed zone is very sensitive to the reproducibility of the assay → Robustness of the methodology to be evaluated**



Bibliography

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