

Dry-off treatment of dairy-cows

Methods to guide targeted antimicrobial use

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Objectives

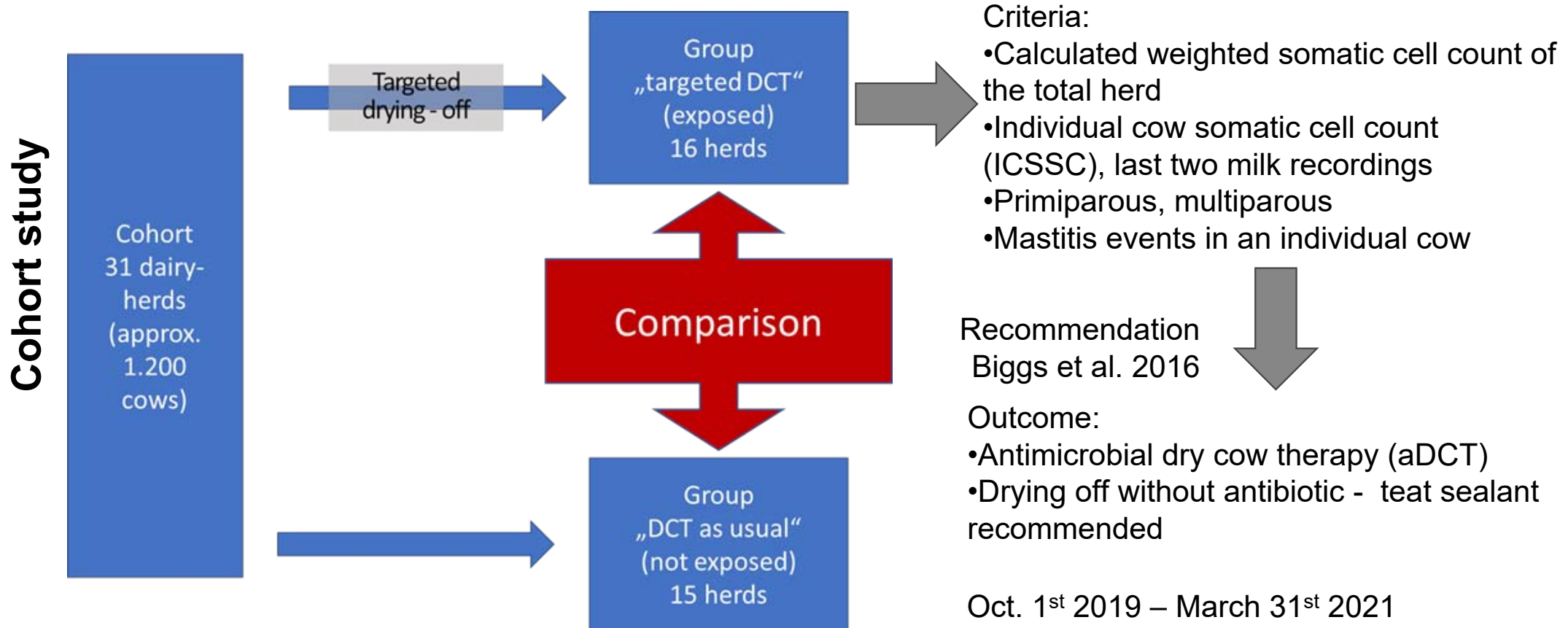
Development of a personalised **drying-off strategy** for **individual dairy farms**.

Minimisation of antibiotic use without negatively affecting udder health.

Development of targeted „**Decision Support Tools**“ to enable implementation of cow-specific drying off strategy.

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Bacteriological culture of milk samples

- a) before drying-off,
- b) after calving,
- c) for every mastitis case

Collation of antimicrobial use data with respect to drying-off

Assessment of the drying-off strategy cure rate, new infections, antimicrobial use

Development of a data-based, cow-specific dry-off recommendation

GEE, random forest (study data set, extended data set)

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Cohort study, CASE herds: targeted antimicrobial dry cow therapy (ADCT)

Weighted SCC of the total herd x 10 ³ cells / ml	ICSCC- threshold x 10 ³ cells / ml	ICSCC- threshold x 10 ³ cells / ml
	Multiparous cows	Primiparous cows
< 100	250	200
100 - 150	200	150
150 - 200	150	100
200 - 250	100	50
> 250	Targeted ADCT may not be appropriate!	

Biggs et al., 2016: Antibiotic dry cow therapy: where next?
 Veterinary Record **178**, 93–94.

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Cohort study – results

- Bacteriological examination of milk samples before drying-off (detection of bacteria by culture)
 - Detection of MAJOR- / MINOR-pathogens
 - Factors associated with the detection of pathogens
- Evaluation of the drying-off strategy
 - Cure rates, new infection rates
 - Antimicrobial use

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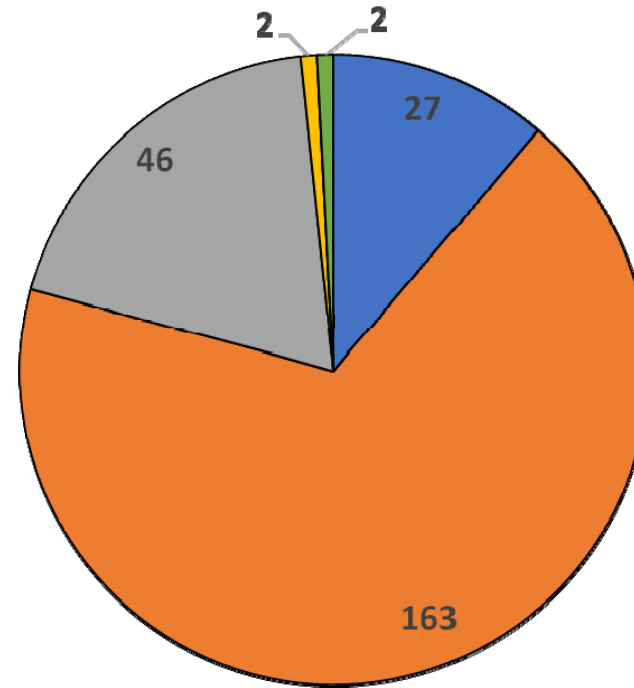
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Detection of bacteria by culture before drying off (quarter milk samples)

Species	N
Staph. aureus	27
Streptococci + Enterococci	163
Enterobacteria	46
Trueperella spp.	2
Mixed infections	2
MINOR pathogens*	260
Negative, contaminated	3,741
	4,241

*Coagulase negative Staphylococci + Corynebacteria

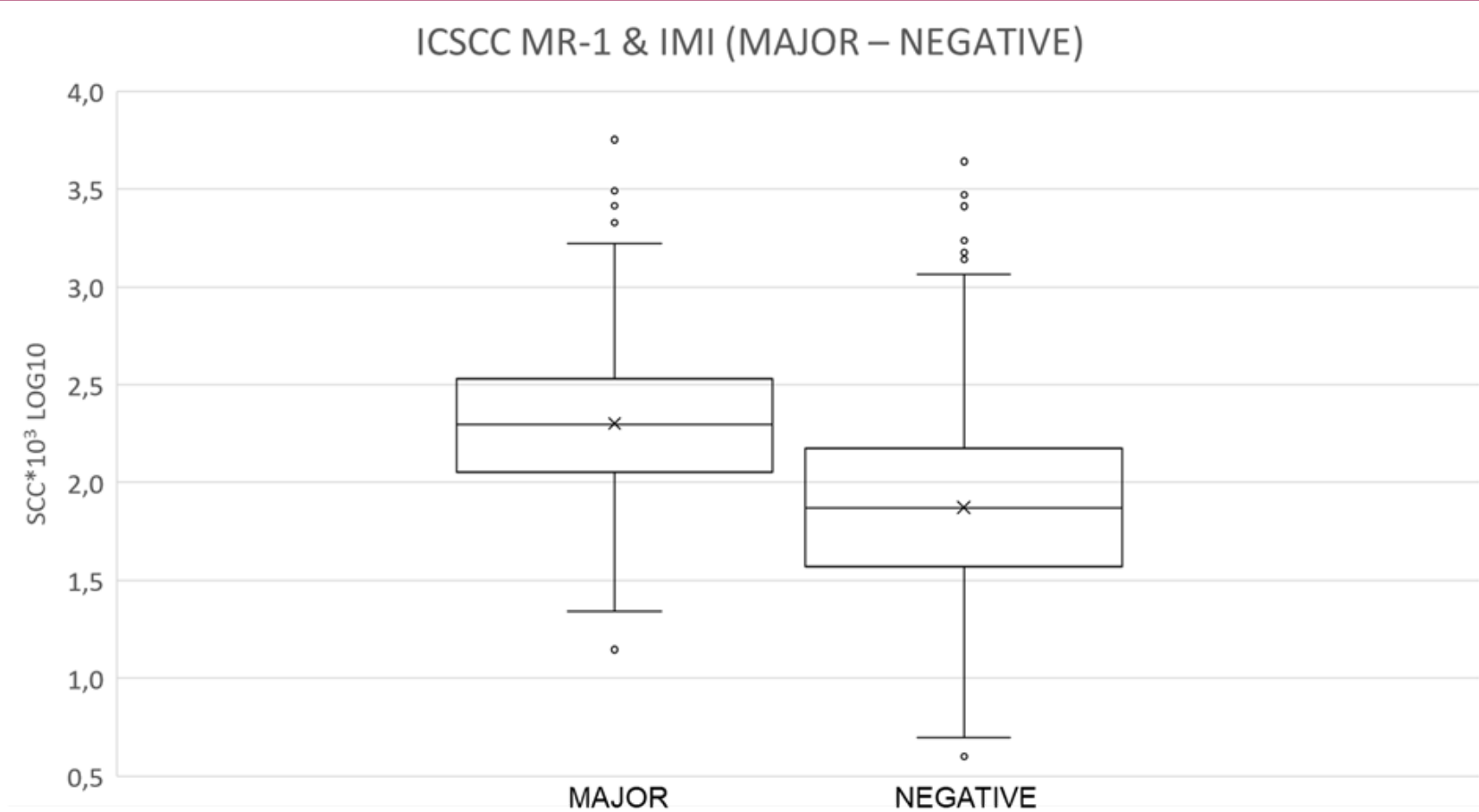


Pathogens	N	%
MAJOR	240	5.7
MINOR	260	6.1
NEGATIVE	3,741	88.2

- Staph. aureus
- Streptococci + Enterococci
- Enterobacteria
- Trueperella
- Yeasts, Algae
- Mixed infections

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Evaluation of the drying-off strategy: cure rates, new infection rates

Status of infection	Total		Dry cow therapy				CASE				CONTROL			
			AM		No AM		AM		no AM		AM		no AM	
No infection	404	58%	215	50%	189*	73%	80	47%	107	75%	135	51%	82	70%
New infection	89	13%	47	11%	42	16%	22	13%	24	17%	25	9%	18	15%
Cure	171	25%	146*	34%	25	10%	58	34%	10	7%	88	33%	15	13%
Persistently infected	30	4%	26	6%	4	2%	10	6%	2	1%	16	6%	2	2%
	694	100%	434	100%	260	100%	170	100%	143	100%	264	100%	117	100%

* significant Pearson's Chi-squared test

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Evaluation of the drying-off strategy: cure rates, new infection rates

			Dry cow therapy				CASE				CONTROL			
Status of infection	total		AM		no AM		AM		No AM		AM		No AM	
No infection	404	58%	215	50%	189*	73%	80	47%	107	75%	135	51%	82	70%
New infection	89	13%	47	11%	42	16%	22	13%	24	17%	25	9%	18	15%
Cure	171	25%	146*	34%	25	10%	58*	34%	10	7%	88*	33%	15	13%
Persistently infected	30	4%	26	6%	4	2%	10	6%	2	1%	16	6%	2	2%
	694	100%	434	100%	260	100%	170	100%	143	100%	264	100%	117	100%

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Evaluation of the drying-off strategy: antimicrobial usage (AM-DCT)

- **Formula to calculate antimicrobial usage for dry-off therapy**

$$\begin{aligned} & \# TD_{365} DCT \\ &= \sum_{i=1}^n \frac{\# UD/udder \text{ (route intramam - DC) in period } P}{\# cow \cdot \text{days in period } P \text{ (days)}} \times 365 \times \frac{\text{calving interval}^1 \text{ in period } P \text{ (herd, days)}}{365} \\ & \times \left(1 + \frac{\# cow LN = 1^2 \cdot \text{days in period } P \text{ (days)}}{\# cow \cdot \text{days in period } P \text{ (days)}} \right) \end{aligned}$$

= number of treatment days / cow / year / for aDCT = number of unit doses (UD) per udder given to any cow of a population within 1 year (1 UD = 4 DCDvet = 4 injectors of an antibiotic licensed for intramammary use in dry-cow therapy).

¹Correction for calving interval

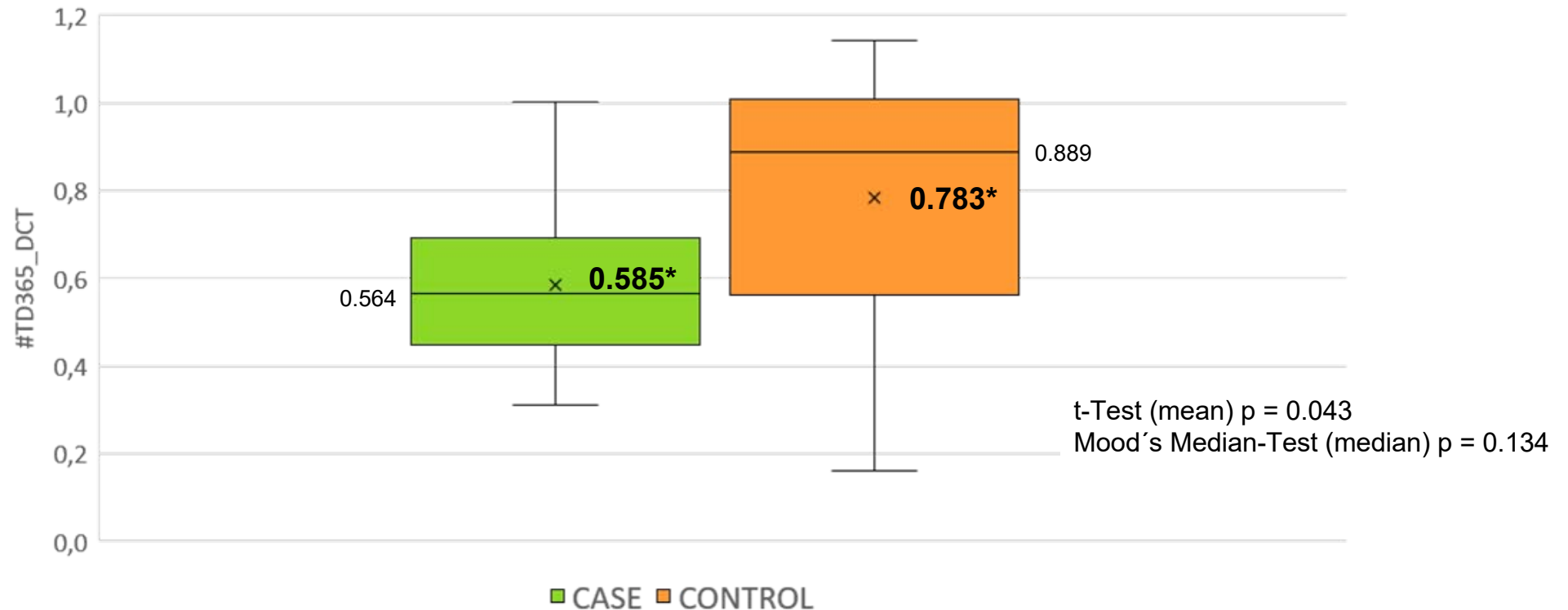
²Correction for herd replacement rate (heifers)

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Evaluation of the drying-off strategy: antimicrobial usage (AM-DCT)

Antimicrobial dry-cow therapy: CASE vs. CONTROL



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- **Statistical modelling (data-based selection method):**

Data collected in the cohort study was used to model the result of bacteriological milk testing at drying off (binary result (major / negative) as target variable).

- Expert-knowledge based method (Biggs et al., 2016)

- ICSCC, SCC-herd, lactation (1, >1)

- Prediction models:

- GEE (Generalized Estimating Equations) model

- Random forest model (Breiman, 2001)

- ICSCC, SCC-herd, lactation, mastitis diagnosis, milk performance recordings, recordings from health monitoring program

Comparison of the two data-based approaches with the expert-knowledge based method

- Field study recommendations were evaluated for the same test set

- Performance measures: accuracy, sensitivity, specificity, and f1-Score

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Comparison of predictions for a positive result of bacteriological milk tests before drying off for the D4Dairy data set (121 observations)

	Recommended ADCT (Biggs et al. 2016)	Generalized Estimating Equations*	Random Forest
Accuracy	0.752	0.719	0.876
Sensitivity	0.652	0.739	0.565
Specificity	0.776	0.714	0.949
F1-Score	0.500	0.500	0.634

* A cut-off value of 0.19 was used to classify the GEE prediction into positive bacteriological milk cultures or negative tests. This was determined using an ROC analysis.

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Comparison of predictions for a positive result of bacteriological milk tests before drying off for the extended data set (2 838 observations)

	Recommended ADCT (Biggs et al. 2016)	Random Forest
Accuracy	0.645	0.700
Sensitivity	0.593	0.519
Specificity	0.681	0.827
F1-Score	0.579	0.588

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- **Summary:**

- Evaluation of the drying-off strategy - cure rates, new infection rates
 - New infections: no significant difference between AM-DCT and no-DCT
 - Cure, new infections: no significant differences between CASE and CONTROL herds
- A selective dry cow therapy recommendation led to a reduced mean antimicrobial usage (AM-DCT)
 - Significant (weak, p-value = 0.043) mean difference between CASE herds (assisted targeted DCT) and CONTROL herds (selective DCT – blanket AM-DCT)
- Data based selection method
 - Prediction models demonstrate, that selection for ADCT based on cell count thresholds alone recommends the use of antibiotics more often than is necessary
 - Relationship between udder infections, milk performance test results, lactation age and herd health indicators is complex and could not be well explained with a GEE model
 - With a statistical prediction model (random forest), an even more precise selection for ADCT could be made
 - Due to the marginally lower sensitivity, a few more infections would have been missed compared to the SDCT method



Thank you for your attention!



D4Dairy

Research partners:



Company partners:



Cooperation partners:



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