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How and why to monitor social networks in dairy cows

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1 Introduction

One of the most important environmental influences on a cow is the social environment induced by surrounding cows. Living in groups leads to affiliative and agonistic interactions between the individuals of a herd. Affiliative and aggressive interactions create balance and structure the herd (Tucker, 2017). Affiliative interactions, such as allogrooming and proximity, contribute to forming and maintaining positive social relationships within the herd (Val-Laillet et al., 2009). In contrast, negative interactions such as aggressive contacts, displacements or replacements, motivated by access to resources and housing conditions, among other factors, contribute to establishing dominance relationships between the individuals (Foris et al., 2021).

These relationships can be affected by daily husbandry procedures on dairy farms, such as re-grouping of animals according to age or production stage or insufficient space allowance (Bouissou et al., 2001; Rocha et al., 2020). Frequent regrouping challenges dominance relationships and social preferences, which can increase levels of aggression within the herd (Raussi et al., 2005; Schirmann et al., 2011). Thus, daily husbandry practices on dairy farms can unintentionally disrupt these social interactions, impacting animal welfare and production (Bouissou et al., 2001; Thompson et al., 2022; von Keyserlingk et al., 2009). Moreover, social interactions play a role in the transmission of contagious diseases (Burke et al., 2022; de Freslon et al., 2019). These studies described the relationship between social network structure and the incidence of respiratory and reproductive diseases in cattle.

Due to the importance of social behavior in cattle's daily activities, scientific research has focused on understanding their social interactions. Visual observations have been the main tool to study social interactions in cattle (Sahu et al., 2020). Visual observations are a robust

and valid method based on human assessments and decisions. However, they are sensitive to inter-observer variability, laborintensive and generally typically limited by the duration of the observation and the number of animals that can be monitored simultaneously. Precision livestock farming (PLF) has opened new possibilities for monitoring and studying behavior in cattle. Real-time location systems (RTLS), for example, allow the continuous monitoring of area utilization (Ternman et al., 2019; Tullo et al., 2016) and proximity between individuals (Ben-Meir et al., 2023; Boyland et al., 2016; Chopra et al., 2020; Rocha et al., 2020).

One of the main goals of the research program on cow social interaction and disease transmission (CSI: DT) was to develop data analysis tools for understanding social contact information to be used to minimize disease transmission within dairy farms. We summarize our publications (Table 1) and experiences from this research program into two opinion papers and give advice for future research within the field. Here, we focus on approaches to efficiently monitor and analyze social network information in dairy cattle and its future implications.

2 Social interactions

2.1 Monitoring social contacts

An RTLS based on ultra-wideband (UWB) technology works with tags attached to the cows which communicate with anchors placed in the barn's ceiling. The positioning of a cow is determined by the tag's connection to multiple anchors. Conducting in-depth analyses of animal movement and social contacts requires understanding the data quality by assessing the noise and accuracy of the position, where data is lost, and how accurately missing positions can be interpolated.

In our studies, we have collected positioning data using the CowView system developed by GEA, which guarantees an error distance of approximately 0.50 m, with a reported accuracy of 16 cm (Meunier et al., 2018). A validation study conducted in both a Swedish and a Dutch farm revealed a mean error distance of the positioning data of 0.78 m and 0.54 m for 13 and 21 fixed tags installed in each farm, respectively (Hansson et al., 2023). The quality of the positioning data was best in the center of the farm where the tags on the cows can easily connect to many anchors in the ceiling (Ren et al., 2021b). The system achieves a 95% accuracy in detecting zone-related behavioral activities based on the floorplans of the barn (Tullo et al., 2016). Ren et al. (2021b) found that missing data averaged 31.29% (~7.5 h/d) and 19.97% (~4.8 h/d) of the day for the Swedish and Dutch farm, with the most common scenario being a single second missing. Interpolating missing position data with the Modified Akima algorithm achieved an average interpolation accuracy of 0.17 m (Ren et al., 2022).

Data collected using UWB systems seem satisfactory for detecting proximity between cows. Nevertheless, we removed proximity interactions that lasted less than 10 minutes over an entire day in our analysis to avoid considering stochastic short contacts due to the limited space of the barn. We attempted to classify affiliative and agonistic interactions by detecting displacements at the feeding tables using the UWB data and video recording but were unsuccessful. The main challenge was that the accuracy of the UWB system was insufficient to detect cow displacements in the open structure feeding bunks. Additionally, the CowView system's internal Kalman filter could delay the detection of animal movements, distorting the actual trajectory.

2.2 Analyzing social contacts

The benefits of using RTLS for studying social contacts in livestock are evident, as it allows the continuous monitoring of proximity between individuals in the herd over long periods. However, proximity contacts do not imply social or physical interaction between individuals. Therefore, their applicability has been criticized in comparison to visual observational studies, which allow distinguishing between different types of social contacts. Nevertheless, preferential relationships in dairy cattle are expressed through social grooming or by spending more time in proximity (Val-Laillet et al., 2009). Previous studies have described a positive correlation between affiliative interactions and longlasting proximity contacts between dairy cows and calves, whereas short encounters were associated with agonistic interactions in calves (Ben-Meir et al., 2023; Boyland et al., 2016).

Scientific studies based on RTLS have described findings consistent with previous literature based on visual observations (e.g. Rocha et al., 2020; Hansson et al., 2023; Boyland et al., 2016), supporting the adoption of this technology to comprehend social behavior in dairy cattle. However, this approach requires some assumptions, such as time, distance and frequency thresholds. Social contacts have been investigated by using distance and cumulative interaction time information (Hansson et al., 2023; Marina et al., 2024c; Rocha et al., 2020) or distance and number of contacts (Boyland et al., 2016; Chopra et al., 2020; Raussi et al., 2010; Vázquez-Diosdado et al., 2023). These studies individually validated the proximity thresholds adopted by varying the temporal and spatial thresholds, supporting the robustness of using proximity to quantify social contacts.

Pairwise interaction data can be used to construct social networks. This information is usually modeled by applying individual- or relational-level models (O'Malley and Marsden, 2008). Hansson et al. (2023) applied individual-level models to explore the relationship between different cow's characteristics and the contact rate on social networks. Applying these models allowed the authors to define the association between parity and days in milk with the number of contacts a cow establishes daily with its herd mates. Marina et al. (2024c) applied a relational-level model known as the separable temporal exponential random graph model described by Krivitsky and Goodreau (2016), to explore longitudinal networks and study the formation and persistence of social contacts in dairy cattle. Implementing this longitudinal relational-level model allowed the authors to describe the kindergarten effect: cows born within seven days of each other showed more persistent contact with each other than with the rest

Торіс	Title	Year	Authors	DOI
Data quality	Where do we find missing data in a commercial real-time location system? Evidence from 2 dairy farms	2021	Keni Ren, Per Peetz Nielsen, Moudud Alam and Lars Rönnegård.	10.3168/JDSC.2020-0064
Data quality	Interpolation methods to improve data quality of indoor positioning data for dairy cattle	2022	Keni Ren, Moudud Alam, Per Peetz Nielsen, Maya Gussmann and Lars Rönnegård.	10.3389/ FANIM.2022.896666
Social contacts	Parity and days in milk affect cubicle occupancy in dairy cows	2021	Mikhail Churakov, Anna Maria Silvera, Maya Gussmann and Per Peetz Nielsen.	10.1016/ J.APPLANIM.2021.105494
Social contacts	Cow characteristics associated with the variation in number of contacts between dairy cows	2023	Ida Hansson, Anna Silvera, Keni Ren, Svenja Woudstra, Anna Skarin, Freddy Willem Fikse, Per Peetz Nielsen and Lars Rönnegård.	10.3168/JDS.2022-21915
Social contacts	New insight into social relationships in dairy cows, and how time of birth, parity and relatedness affect spatial interactions later in life	2023	Hector Marina, Keni Ren, Ida Hansson, Freddy Willem Fikse, Per Peetz Nielsen and Lars Rönnegård.	10.3168/JDS.2023-23483
Social contacts	Social network analysis to predict social behavior in dairy cattle	2023	Hector Marina, Freddy Willem Fikse and Lars Rönnegård.	10.3168/JDSC.2023-0507
Social contacts	Multiple factors shape social contacts in dairy cows	2024	Hector Marina, Per Peetz Nielsen, Freddy Willem Fikse and Lars Rönnegård.	10.1016/ J.APPLANIM.2024.106366
Milking behavior	Associations of parity and lactation stage with the order cows enter the milking parlor	2023	Ida Hansson and Svenja Woudstra.	10.3168/JDSC.2023-0491
Infectious pathogens	Strain diversity and infection durations of Staphylococcus spp. and Streptococcus spp. causing intramammary infections in dairy cows	2023	Svenja Woudstra, Nicole Wente, Yanchao Zhang, Stefanie Leimbach, Maya Katrin Gussmann, Carsten Kirkeby and Volker Krömker.	10.3168/JDS.2022-22942
Infectious pathogens	Reservoirs of Corynebacterium spp. in the environment of dairy cows	2023	Svenja Woudstra, Anneke Lücken, Nicole Wente, Yanchao Zhang, Stefanie Leimbach, Maya Katrin Gussmann, Carsten Kirkeby and Volker Krömker.	10.3390/ PATHOGENS12010139
Infectious pathogens	Reservoirs of Staphylococcus spp. and Streptococcus spp. associated with intramammary infections of dairy cows	2023	Svenja Woudstra, Nicole Wente, Yanchao Zhang, Stefanie Leimbach, Carsten Kirkeby, Maya Katrin Gussmann and Volker Krömker.	10.3390/ Pathogens12050699

TABLE 1 Scientific publications published during the project.

Scientific publications published during the lifetime of the project (2020/01/01 - 2024/12/31)

of the herd. Considering these factors together with the internal structure of social networks is crucial to understanding the occurrence and persistence of social contacts in dairy cattle (Marina et al., 2024a).

RTLS data can also be used to discover cubicle occupancy patterns of dairy cows in the barn. Churakov et al. (2021) used the aggregated positioning data within the CowView system with predicted activity (e.g., walking, eating, in-cubicle). They found that cubicle occupancy patterns are related to parity and lactation stage and that cows choose cubicles close to individuals with similar attributes. High parity cows used the cubicles close to the milking area more frequently, while first lactation cows occupied less busy barn areas. In addition, Marina et al. (2024b) demonstrated the importance of considering the spatial preferences of individuals when studying social interactions in dairy cattle.

3 Genetics

Only a few studies have investigated the genetic component of sociability in dairy cattle (e.g. Gutiérrez-Gil et al., 2008; Marina et al., 2024c; Rönnegård et al., 2022; Wang et al., 2023). Hansson et al. (2023) investigated the repeatability of social behavior in dairy cows over consecutive days by estimating the proportion of variation in contact rate between individuals relative to the total variance, and determined that it was approximately 30%. Rönnegård et al. (2022) used these data to investigate how much of the variation between animals was due to genetics and estimated heritabilities (proportion of genetic variance relative to total variance) of 0 to 0.20, albeit with large standard errors. Similar heritability estimates were reported in *Bos indicus* for other social measures (dominance value and social hierarchy), with values of 0.23 and 0.25, respectively (de Paula Soares Valente et al., 2023). These estimates correspond well to heritability estimates of similar traits in pigs (Agha et al., 2022a, 2022b).

Not only is the number of contacts a cow has heritable, but the effect a cow has on other cows' milk production might be heritable too. This is known as indirect genetic effects (IGEs) (Moore et al., 1997). IGEs can make a significant contribution to heritable variation in livestock species, helping to reduce disease transmission, stress levels and overall herd health issues (Peeters et al., 2012), for instance. There are several examples of the benefits of understanding IGEs in different livestock species, such as the cannibalistic behavior in laying hens (Alemu et al., 2016), tail biting in pigs (Camerlink et al., 2015), and the outcome of dyadic contacts in deer (Wilson et al., 2011). Using methods that target both the direct and indirect genetic effects for genetic selection, is a

promising tool to simultaneously improve production and welfare through the selection programs in dairy cattle (Ellen et al., 2014).

4 Discussion

The continuous collection of proximity and area utilization data for all individuals over time has allowed us to gain new insights into the behavior of dairy cattle in free-stall barns. The UWB data have facilitated the description of how cubicle occupancy varies within and between lactations, revealed the heterogeneity of social networks in dairy cattle and how long-lasting preferences for social contacts between cows continue even between lactations. Similar technologies could have been used to obtain this information, such as Bluetooth wireless technology or proximity loggers. However, UWB systems provide fairly accurate positions compared to Bluetooth (see Huhtala et al., 2007; Bloch and Pastell, 2020). Spatial proximity loggers could also provide temporal proximity information, although the area of the barn where these interactions occurred will be challenging to determine.

The findings from this project could have benefited from combining UWB data with video information and data from proximity loggers to overcome the limitations of the system and increase the accuracy of social networks. The integration of UWB data with video image analysis could provide specific information about certain behaviors and allow the detection of the affiliative and agonistic social interactions (McDonagh et al., 2021; Meunier et al., 2018; Ren et al., 2021a). In addition, the fusion of indoor and outdoor sensors would allow continuous monitoring of all individuals throughout the year also in production systems with pasture access, enhancing the use of this information to improve farm management practices and gain valuable information on animal behavior. Altogether, RTLS data is a valuable tool that could contribute to understanding the social behavior of dairy cattle, optimize farm management practices, and improve animal health, production and welfare of dairy cattle.

Author contributions

HM: Writing – original draft, Writing – review & editing. IH: Writing – original draft, Writing – review & editing. KR: Writing – original draft, Writing – review & editing. FF: Writing – review & editing. MG: Writing – review & editing. PN: Writing – review &

References

Agha, S., Foister, S., Roehe, R., Turner, S. P., and Doeschl-Wilson, A. (2022a). Genetic analysis of novel behaviour traits in pigs derived from social network analysis. *Genes (Basel)* 13, 561. doi: 10.3390/GENES13040561/S1

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Alemu, S. W., Calus, M. P. L., Muir, W. M., Peeters, K., Vereijken, A., and Bijma, P. (2016). Genomic prediction of survival time in a population of brown laying hens showing cannibalistic behavior. *Genet. Sel. Evol* 48, 1–10. doi: 10.1186/S12711-016-0247-4/TABLES/7

Ben-Meir, Y. A., Garcia, F., Cohen-Zinder, M., and Shabtay, A. (2023). Use of proximity loggers to Estimate Affiliative and Agonistic Relationships among Group-housed Holstein calves. J. Appl. Anim. Welf. Sci. 28, 209–221. doi: 10.1080/10888705.2023.2250262

Agha, S., Turner, S. P., Lewis, C. R. G., Desire, S., Roehe, R., and Doeschl-Wilson, A. (2022b). Genetic associations of novel behaviour traits derived from social network analysis with growth, feed efficiency, and carcass characteristics in pigs. *Genes (Basel)* 13, 1616. doi: 10.3390/GENES13091616/S1

Bloch, V., and Pastell, M. (2020). Monitoring of cow location in a barn by an opensource, low-cost, low-energy bluetooth tag system. *Sensors* 20, 3841. doi: 10.3390/ S20143841

Bouissou, M.-F., Boissy, A., Le Neindre, P., and Veissier, I. (2001). The social behaviour of cattle, in: Social Behaviour in Farm Animals. *CABI Publishing* pp, 113–145. doi: 10.1079/9780851993973.0113

Boyland, N. K., Mlynski, D. T., James, R., Brent, L. J. N., and Croft, D. P. (2016). The social network structure of a dynamic group of dairy cows: From individual to group level patterns. *Appl. Anim. Behav. Sci* 174, 1-10. doi: 10.1016/J.APPLANIM.2015.11.016

Burke, K. C., do-Nascimento-Emond, S., Hixson, C. L., and Miller-Cushon, E. K. (2022). Social networks respond to a disease challenge in calves. *Sci. Rep* 121 12, 1–10. doi: 10.1038/s41598-022-13088-2

Camerlink, I., Ursinus, W. W., Bijma, P., Kemp, B., and Bolhuis, J. E. (2015). Indirect genetic effects for growth rate in domestic pigs alter aggressive and manipulative biting behaviour. *Behav. Genet* 45, 117. doi: 10.1007/S10519-014-9671-9

Chopra, K., Hodges, H. R., Barker, Z. E., Vázquez Diosdado, J. A., Amory, J. R., Cameron, T. C., et al. (2020). Proximity interactions in a permanently housed dairy herd: network structure, consistency, and individual differences. *Front. Vet. Sci* 7. doi: 10.3389/FVETS.2020.583715/BIBTEX

Churakov, M., Silvera, A. M., Gussmann, M., and Nielsen, P. P. (2021). Parity and days in milk affect cubicle occupancy in dairy cows. *Appl. Anim. Behav. Sci* 244, 105494. doi: 10.1016/J.APPLANIM.2021.105494

de Freslon, I., Martínez-López, B., Belkhiria, J., Strappini, A., and Monti, G. (2019). Use of social network analysis to improve the understanding of social behaviour in dairy cattle and its impact on disease transmission. *Appl. Anim. Behav. Sci* 213, 47–54. doi: 10.1016/J.APPLANIM.2019.01.006

de Paula Soares Valente, J., Tenffen-De-Sousa, K., Deniz, M., Figueiredo Martins Bonilha, S., Eugènia Zerlotti Mercadante, M., and Talarico Dias, L. (2023). Heritability and genetic association of social organization traits with feeding behavior, feed efficiency and growth in Bos indicus. *Appl. Anim. Behav. Sci.* 265, 105976. doi: 10.1016/j.applanim.2023.105976

Ellen, E. D., Bas Rodenburg, T., Albers, G. A. A., Elizabeth Bolhuis, J., Camerlink, I., Duijvesteijn, N., et al. (2014). The prospects of selection for social genetic effects to improve welfare and productivity in livestock. *Front. Genet* 5. doi: 10.3389/ FGENE.2014.00377/ABSTRACT

Foris, B., Lecorps, B., Krahn, J., Weary, D. M., and von Keyserlingk, M. A. G. (2021). The effects of cow dominance on the use of a mechanical brush. *Sci. Rep* 111 11, 1–7. doi: 10.1038/s41598-021-02283-2

Gutiérrez-Gil, B., Ball, N., Burton, D., Haskell, M., Williams, J. L., and Wiener, P. (2008). Identification of quantitative trait loci affecting cattle temperament. *J. Hered* 99, 629–638. doi: 10.1093/JHERED/ESN060

Hansson, I., Silvera, A., Ren, K., Woudstra, S., Skarin, A., Fikse, W. F., et al. (2023). Cow characteristics associated with the variation in number of contacts between dairy cows. J. Dairy Sci 106, 2685–2699. doi: 10.3168/JDS.2022-21915

Huhtala, A., Suhonen, K., Mäkelä, P., Hakojärvi, M., and Ahokas, J. (2007). Evaluation of instrumentation for cow positioning and tracking indoors. *Biosystems Engr.* 96, 399–405. doi: 10.1016/j.biosystemseng.2006.11.013

Krivitsky, P. N., and Handcock, M. S. (2014). A Separable Model for Dynamic Networks. J. R. Stat. Soc. Ser. B Stat. Methodol. 76, 29–46. doi: 10.1111/RSSB.12014

Marina, H., Fikse, W. F., and Rönnegård, L. (2024a). Social network analysis to predict social behavior in dairy cattle. *JDS Commun.* 5, 608–612. doi: 10.3168/JDSC.2023-0507

Marina, H., Nielsen, P. P., Fikse, W. F., and Rönnegård, L. (2024b). Multiple factors shape social contacts in dairy cows. *Appl. Anim. Behav. Sci* 278, 106366. doi: 10.1016/ J.APPLANIM.2024.106366

Marina, H., Ren, K., Hansson, I., Fikse, F., Nielsen, P. P., and Rönnegård, L. (2024c). New insight into social relationships in dairy cows and how time of birth, parity, and relatedness affect spatial interactions later in life. *J. Dairy Sci* 107, 1110–1123. doi: 10.3168/JDS.2023-23483

McDonagh, J., Tzimiropoulos, G., Slinger, K. R., Huggett, Z. J., Bell, M. J., and Down, P. M. (2021). Detecting dairy cow behavior using vision technology. *Agric* 11, 675. doi: 10.3390/AGRICULTURE11070675

Meunier, B., Pradel, P., Sloth, K. H., Cirié, C., Delval, E., Mialon, M. M., et al. (2018). Image analysis to refine measurements of dairy cow behaviour from a real-time location system. *Biosyst. Eng* 173, 32–44. doi: 10.1016/J.BIOSYSTEMSENG.2017.08.019

Moore, A. J., Brodie, E. D., and Wolf, J. B. (1997). Interacting phenotypes and the evolutionary process: I. direct and indirect genetic effects of social interactions. *Evol.* (*N. Y*) 51, 1352–1362. doi: 10.1111/J.1558-5646.1997.TB01458.X

O'Malley, A. J., and Marsden, P. V. (2008). The analysis of social networks. *Health Serv. Outcomes Res. Methodol* 8, 222. doi: 10.1007/S10742-008-0041-Z

Peeters, K., Eppink, T. T., Ellen, E. D., Visscher, J., and Bijma, P. (2012). Indirect genetic effects for survival in domestic chickens (Gallus gallus) are Magnified in crossbred genotypes and show a parent-of-origin effect. *Genetics* 192, 705–713. doi: 10.1534/GENETICS.112.142554/-/DC1

Raussi, S., Boissy, A., Delval, E., Pradel, P., Kaihilahti, J., and Veissier, I. (2005). Does repeated regrouping alter the social behaviour of heifers? *Appl. Anim. Behav. Sci* 93, 1–12. doi: 10.1016/J.APPLANIM.2004.12.001

Raussi, S., Niskanen, S., Siivonen, J., Hänninen, L., Hepola, H., Jauhiainen, L., et al. (2010). The formation of preferential relationships at early age in cattle. *Behav. Processes* 84, 726–731. doi: 10.1016/J.BEPROC.2010.05.005

Ren, K., Alam, M., Nielsen, P. P., Gussmann, M., and Rönnegård, L. (2022). Interpolation methods to improve data quality of indoor positioning data for dairy cattle. *Front. Anim. Sci* 0. doi: 10.3389/FANIM.2022.896666

Ren, K., Bernes, G., Hetta, M., and Karlsson, J. (2021a). Tracking and analysing social interactions in dairy cattle with real-time locating system and machine learning. *J. Syst. Archit* 116, 102139. doi: 10.1016/J.SYSARC.2021.102139

Ren, K., Nielsen, P. P., Alam, M., and Rönnegård, L. (2021b). Where do we find missing data in a commercial real-time location system? Evidence from 2 dairy farms. *JDS Commun* 2, 345–350. doi: 10.3168/JDSC.2020-0064

Rocha, L. E. C., Terenius, O., Veissier, I., Meunier, B., and Nielsen, P. P. (2020). Persistence of sociality in group dynamics of dairy cattle. *Appl. Anim. Behav. Sci* 223, 104921. doi: 10.1016/J.APPLANIM.2019.104921

Rönnegård, L., Hansson, I., and Fikse, W. F. (2022). Heritability of social interactions in dairy cattle, in: Proceedings of 12th World Congress on Genetics Applied to Livestock Production (WCGALP). *Wageningen Acad. Publishers* pp, 486–489. doi: 10.3920/978-90-8686-940-4_109

Sahu, B. K., Parganiha, A., and Pati, A. K. (2020). Behavior and foraging ecology of cattle: A review. J. Vet. Behav 40, 50-74. doi: 10.1016/j.jveb.2020.08.004

Schirmann, K., Chapinal, N., Weary, D. M., Heuwieser, W., and von Keyserlingk, M. A. G. (2011). Short-term effects of regrouping on behavior of prepartum dairy cows. *J. Dairy Sci* 94, 2312–2319. doi: 10.3168/JDS.2010-3639

Ternman, E., Nilsson, E., Nielsen, P. P., Pastell, M., Hänninen, L., and Agenäs, S. (2019). Rapid eye movement sleep time in dairy cows changes during the lactation cycle. J. Dairy Sci 102, 5458–5465. doi: 10.3168/JDS.2018-15950

Thompson, J. S., Hudson, C. D., Huxley, J. N., Kaler, J., Robinson, R. S., Woad, K. J., et al. (2022). A randomised controlled trial to evaluate the impact of indoor living space on dairy cow production, reproduction and behaviour. *Sci. Rep* 121 12, 1–21. doi: 10.1038/s41598-022-07826-9

Tucker, C. B. (2017). Behaviour of cattle, in: The Ethology of Domestic Animals, 3rd Edition: An Introductory Text. *CAB Int* pp, 189–198. doi: 10.1079/9781786391650.0189

Tullo, E., Fontana, I., Gottardo, D., Sloth, K. H., and Guarino, M. (2016). Technical note: Validation of a commercial system for the continuous and automated monitoring of dairy cow activity. *J. Dairy Sci* 99, 7489–7494. doi: 10.3168/JDS.2016-11014

Val-Laillet, D., Guesdon, V., von Keyserlingk, M. A. G., de Passillé, A. M., and Rushen, J. (2009). Allogrooming in cattle: Relationships between social preferences, feeding displacements and social dominance. *Appl. Anim. Behav. Sci* 116, 141–149. doi: 10.1016/J.APPLANIM.2008.08.005

Vázquez-Diosdado, J. A., Occhiuto, F., Carslake, C., and Kaler, J. (2023). Familiarity, age, weaning and health status impact social proximity networks in dairy calves. *Sci. Rep* 131 13, 1–10. doi: 10.1038/s41598-023-29309-1

von Keyserlingk, M. A. G., Rushen, J., de Passillé, A. M., and Weary, D. M. (2009). Invited review: The welfare of dairy cattle—Key concepts and the role of science. J. Dairy Sci 92, 4101-4111. doi: 10.3168/JDS.2009-2326

Wang, Z., Doekes, H., and Bijma, P. (2023). Towards genetic improvement of social behaviours in livestock using large-scale sensor data: data simulation and genetic analysis. *Genet. Sel. Evol* 55, 1–14. doi: 10.1186/S12711-023-00840-Z/TABLES/5

Wilson, A. J., Morrissey, M. B., Adams, M. J., Walling, C. A., Guinness, F. E., Pemberton, J. M., et al. (2011). Indirect genetics effects and evolutionary constraint: an analysis of social dominance in red deer, Cervus elaphus. J. Evol. Biol 24, 772–783. doi: 10.1111/J.1420-9101.2010.02212.X