

DEVELOPMENT AND VALIDATION OF RISK PREDICTION TOOLS FOR PRESSURE INJURY OCCURRENCE: AN UMBRELLA REVIEW

1. Background

- **Epidemiology and Costs:** Pressure injuries (PIs) have a global prevalence of 13%, with two-thirds being hospital-acquired (HAPI). They impose significant financial burdens, costing \$27 billion annually in the U.S. due to 2.5 million cases, and increase hospital stays by an average of 10 days for patients over 75 years old.
- **Risk Factors and Prevention:** PIs typically develop on bony areas such as heels and coccyx predominantly in patients with limited mobility. Effective prevention involves multifaceted strategies including support surfaces, nutritional supplements, and regular repositioning.
- **Challenges in Risk Assessment:** Traditional risk assessment relies on clinical scales like Braden and Norton, which are subjective. Recent advancements in machine learning and statistical models promise better accuracy but require robust validation.
- **Research Gap:** Systematic reviews on PI risk prediction tools are often limited in scope and vary in quality, complicating comparisons and understanding.
- **Study Objective:** This study conducts an umbrella review to systematically evaluate and summarize evidence on the development, validation, and utility of risk prediction tools for PIs.

2. Methods

- **Protocol and Reporting Standards:**
 - Utilized Cochrane Handbook for umbrellas reviews.
 - Reported findings adhering to PRISMA guidelines.
 - Protocol registered on the Open Science Framework.
- **Literature Search:**
 - Conducted by an experienced information specialist in January 2023.
 - Searches executed in MEDLINE, Embase via Ovid, CINAHL Plus EBSCO, EPISTEMONIKOS, and Google Scholar.
 - Combined systematic review and prognostic search filters with PI-related terms.
 - Independent dual-review of titles, abstracts, and full texts.
- **Eligibility Criteria:**
 - English-language systematic reviews focusing on risk prediction models for adult patients at risk of PI.
 - Included reviews of both clinical tools and models developed using statistical or ML methods, with or without validation.
- **Data Extraction and Quality Assessment:**
 - Utilized CHARMS checklist and Cochrane Prognosis group template for data extraction.
 - AMSTAR-2 adapted for risk prediction models used for assessing methodological quality.
 - Dual-review process for extraction and quality assessment.
- **Synthesis Methods:**
 - Grouped reviews based on reporting of model development and validation.
 - Narrative synthesis and tabulation of methods and performance measures.
 - No statistical synthesis conducted; findings on prognostic accuracy and clinical effectiveness reported separately.

3. Results

- **Characteristics of Included Reviews**
- **Search and Selection Process:**
 - After de-duplication, 6,301 unique records were identified.
 - 110 records assessed in full-text, with 103 obtained and one additional from references.
 - 45 reviews listed available risk prediction models; 28 met eligibility for inclusion.
 - 5 reviews detailed model development and internal validation, one also included external validation. 16 reviews focused on reporting accuracy data, 10 reviews evaluated clinical effectiveness, with three covering both accuracy and effectiveness.

3. Results

- **Review Characteristics:**
 - Publication years ranged from 2006 to 2023.
 - Half of the reviews conducted meta-analyses.
 - Reviews included between 1 and 70 primary studies, with participant numbers ranging from 528 to over 1.27 million.
 - Maximum number of prediction tools reviewed was 35.
- **Inclusion Specifics:**
 - 15 reviews had specific inclusion criteria based on setting or patient population (e.g., hospital inpatients, acute care, surgical patients, long-term care, elderly, bedridden).
 - 8 reviews exclusively analyzed the Braden or Waterlow scales.

Review characteristics	Reviews on model development and validation (N=5) 2014 (2011 – 2023)	Reviews on accuracy or clinical effectiveness (N=23) 2016 (2008 – 2022)
Median (range) year of publication		
Eligibility criteria		
Population A	2 (40)	3 (13)
Any population	1 (20)	0 (0)
Bedridden	1 (20)	1 (22)
Inpatients	0 (0)	12 (52)
Adults	0 (0)	6 (26)
No PIs at baseline	1 (20)	5 (22)
NS		
Setting A	0 (0)	3 (13)
Any healthcare setting	1 (20)	5 (22)
Hospital	0 (0)	2 (9)
Long-term care	0 (0)	6 (26)
Acute care (incl. surgical and ICU)	0 (0)	1 (4)
Long-term or acute	4 (80)	8 (35)
NS		
Prediction models	4 (80)	1 (4)
ML models	1 (20)	0 (0)
ML or statistical models	0 (0)	9 (39)
Any tools/scales	0 (0)	10 (43)
Specified clinical scales	0 (0)	1 (4)
Other	0 (0)	2 (9)
NS		
PI classification system	0 (0)	1 (4)
Any	0 (0)	2 (9)
Accepted standard classifications	0 (0)	3 (13)
Several specified classification systems (NPUAP, EPUAP, AHCPR or TDCPS)	0 (0)	1 (4)
Other	5 (100)	18 (70)
NS		
Source of data	0 (0)	4 (520)
Prospective only	1 (20)	2 (511)
Prospective or retrospective	0 (0)	18 (70)
NS		
Study design restrictions	1 (20)	13 (57)
Yes	0 (0)	4 (17)
No	4 (80)	6 (26)
NS		
Review methods	5 (2 – 8)	6 (2 – 14)
Publication restrictions:		
End date (year)		
2000–2009	0 (0)	3 (13)
2010–2019	1 (20)	16 (70)
2020–2023	4 (80)	4 (17)
Language	4 (80)	9 (39)
English only	0 (0)	2 (9)
2 languages	0 (0)	3 (13)
≥2 languages	0 (0)	4 (17)
No restrictions	1 (20)	5 (22)
NS		
Quality assessment tool A	3 (60)	1 (4)
PROBAST	0 (0)	2 (9)
QUADAS	0 (0)	7 (30)
QUADAS-2	0 (0)	3 (13)
JBI tools	0 (0)	2 (9)
CASP	0 (0)	6 (26)
Other	2 (40)	4 (17)
NS	1 (20)	13 (57)
Meta-analysis included		

Table 1. Summary of included systematic review characteristics

Number of studies (N) of reviews, unless otherwise specified. A review may fall into multiple categories, therefore total number within domain not necessarily equal to 100%. B = bias domains, bibliographies or registries; P = review 54 started before 2000; C = Cochrane; R = randomised; N = not randomised; N = not randomised; P = pressure injury; ICU = intensive care unit; M = machine learning; NPJAP = National Pressure Ulcer Advisory Panel; EPJAP = European Pressure Ulcer Advisory Panel; ACPDR = Agency for Health Care Policy and Research; TDCPR = Tennessee Developmental Classification of Pressure Sores; QJADAS 2.0 = Quality Assessment of Diagnostic Accuracy Studies (version 2); PROBAST = Prediction Model Risk of Bias Assessment; Jo = Joanna Briggs Institute; CASP = Critical Appraisal Skills Programme; FE = fixed effects; RSE = random effects; DA = diagnostic test accuracy.

- **Quality Assessment Tools Used:**
 - PROBAB tool used in four reviews for quality assessment.
 - QUADAS-2 used in seven reviews, and original QUADAS in two.
 - Six reviews did not conduct a quality assessment of primary studies.



Figure 2. Summary of AMSTAR-2 assessment results

3. Results (continued)

Review author (publicatio n year)	DEV/ VAL (no. studies)	Setting of included studies; data sources	Model development algorithms	Internal validation method	Brief description of study quality	Summary of model performance results
Dweekat (2023) (1)	DEV: Unclear	HAPI/CAPIC n=32; SRPI n=2, detection of PI (effect on length of stay) n=1; nursing home residents n=2	LR n=20; RF n=16; DT n=12; SVM n=12; MLP n=2; KNN n=1; LDA n=1; other n=19	CV n=10; split sample n=10; split sample and CV n=8; NS=7	No RoB assessment	N/A
Jiang (2021) (2)	DEV	ICU n=33; operating room n=2; acute care hospital n=1; oncology department n=1; end-of-life care n=1; mobility-related disabilities n=1 EHRs used in all models	DT n=25; SVM n=23; LR n=2; NN n=2; RF n=1; MTS n=1; gradient boosting n=1	RoB assessed using PROBAST. Overall RoB high for all predictive models. All models at high RoB in analysis domain.	Split sample n=4; NS n=9	<i>F</i> -score ranged between 0.377 (ML Su, MTS) and 0.670 (ML Su, LR); g-means ranged between 0.628 (ML Kaepwrag BN) and 0.822 (ML Su MTS); Sensitivity ranged between 0.478 (ML Kwagrag) and 0.848 (ML Yang); Specificity ranged between 0.703 (ML Deng) and 0.988 (ML Su LR)
Ribeiro (2021) (3)	DEV	SRPI cardiovascular n=2; SRPI critical care n=1 EHRs used in n=2 models	ANN n=1; XGBost n=1; RF n=1	Split sample n=2; NS n=1	No RoB assessment	N/A
Shi (2019) (4)	DEV: VAL	General acute care hospital n=7; long-term care n=5; ICU n=4; cardiovascular surgery n=2; trauma and burn centres n=1; rehabilitation units n=1; unclear n=1 Prospective n=10; retrospective n=11 VAL Long-term care (e.g. ICU) n=2; general (acute care) hospital n=2 Prospective n=3; retrospective n=4	LR n=16; cox regression n=5; ANN n=1; DT n=1; decision analysis n=1; C4.5 machine learning (DT) n=1; NS n=1	CV n=1; tree-pruning n=1; split sample n=1; re-sampling n=2; NS n=16	RoB assessed using PROBAST. DEV Overall RoB unclear for two models. Overall RoB high for the remaining 19 models. Analysis and outcome domains were mostly at high RoB. VAL Overall RoB unclear for three validation studies. Overall RoB high for the remaining four validation studies. Analysis and outcome domains were mostly at high RoB.	C-statistic ranged between 0.61 (INTERAI PURS) and 0.90 (TNH-PURP). O/E ratios ranged between 0.91 (Berlowitz MDS) and 1.0 (prePULSE study tool) Pooled C-statistic TNH-PURP: 0.86 (95% CI 0.81–0.90) Fragment scale: 0.79 (95% CI 0.77–0.82), n=2 Berlowitz 11-item model: 0.75 (95% CI 0.74–0.76), n=2 Berlowitz MDS model: 0.73 (95% CI 0.72–0.74), n=2 INTERAI PURS: 0.85 (95% CI 0.60–0.89), n=2 Crompton: 0.81 (95% CI 0.78–0.84), n=2 Pooled O/E ratio 0.90 (95% CI 0.95–1.04), n=2 Berlowitz MDS 0.94 (95% CI 0.88–1.01), n=2
Zhou (2022) (5)	DEV	SRPI n=3; ICU n=1; hospitalised n=2; rehabilitation centre n=1; hospice n=1 EHR n=18; Medical Information Mart for Intensive care II database n=4	LR n=15; RF n=10; DT n=9; SVM n=9; ANN n=8; BN n=3; XGBost n=3; GB n=2; AdaBoost n=1; CONTRIP n=1; LSTM n=1; EN n=1; KNN n=1; MTS n=1; NB	CV n=12; NS n=10	RoB assessed using PROBAST. Overall RoB unclear for five studies. Overall RoB high for 15 models. RoB not assessed in two studies due to use of unstructured data.	F1 kagari ranged between 0.02 (ML Nakagami) and 0.39 (ML Song [2]). AUC ranged between 0.78 (ML Delpanne) and 0.98 (ML Song [2]); sensitivity ranged between 0.08 (ML Cui) and 0.95 (ML Song [2]); specificity ranged between 0.63 (ML Delpanne) and 1 (ML Cui)

TABLE 2. Results of reviews reporting model development and validation

ALC – area under curve; ANN – artificial neural network; BN – Bayesian network; CAPF – community-acquired pneumonia infection; CANPFR – re-current Adverse Cardiovascular Events; CBR – case-based reasoning; CHD – coronary heart disease; EHRs – electronic health records; ELN – elastic net; GR – gradient boosting; HAPY – hospital-acquired pneumonia; ICU – intensive care unit; KNN – k-nearest neighbor; LADG – linear discriminant analysis; LSTM – long short-term memory; LR – logistic regression; ML – machine learning; MLP – multilayer perceptron; MTS – Marikavali-Bagchi Tapsahi system; NB – naïve Bayes; NLP – natural language processing; OR – odds ratio; P – pressure; PR – pressure injury; PROACT – medical Risk of Abs Assessment Tool; RF – random forest; ROB – risk of bias; SRP – surgery-related pressure injury; SVM – support vector machine; VLU – validation.

aAbbreviations from meta-analysis
bAbbreviations from individual studies
cAbbreviations from meta-analyses and individual studies
dAbbreviations from meta-analyses, pooling development and external validation
eAbbreviations from meta-analyses, pooling development and internal validation
fAbbreviations from meta-analyses, pooling development and external validation
gAbbreviations from meta-analyses, pooling development and external validation
hAbbreviations from meta-analyses, pooling development and external validation
iAbbreviations from meta-analyses, pooling development and external validation
jAbbreviations from meta-analyses, pooling development and external validation
kAbbreviations from meta-analyses, pooling development and external validation
lAbbreviations from meta-analyses, pooling development and external validation
mAbbreviations from meta-analyses, pooling development and external validation
nAbbreviations from meta-analyses, pooling development and external validation
oAbbreviations from meta-analyses, pooling development and external validation
pAbbreviations from meta-analyses, pooling development and external validation
qAbbreviations from meta-analyses, pooling development and external validation
rAbbreviations from meta-analyses, pooling development and external validation
sAbbreviations from meta-analyses, pooling development and external validation
tAbbreviations from meta-analyses, pooling development and external validation
uAbbreviations from meta-analyses, pooling development and external validation
vAbbreviations from meta-analyses, pooling development and external validation
wAbbreviations from meta-analyses, pooling development and external validation
xAbbreviations from meta-analyses, pooling development and external validation
yAbbreviations from meta-analyses, pooling development and external validation
zAbbreviations from meta-analyses, pooling development and external validation

23 reviews provided limited details on model development or validation and mainly involved clinically-derived tools such as Braden (n=20) and Waterlow (n=14) scales.

- **Detailed Model Analysis:**
 - Five reviews reported extensive details on model development and validation, covering 62 prediction models, with 40 unique to these reviews.
 - Models predominantly used electronic health records for development, with logistic regression being the most common modeling approach followed by machine learning techniques like random forests and support vector machines.
- **Validation and Quality Assessment:**
 - Internal validation methods were inconsistently reported; some reviews noted a lack of validation information for a significant number of studies.
 - Two reviews utilized PROBAST for quality assessment, finding a high risk of bias in the majority of studies.
- **Model Performance and External Validation:**
 - Performance measures varied widely across reviews, with metrics like C-statistics, F1 scores, and G-means demonstrating considerable variation.
 - External validation was limited but included some models validated in long-term and acute care settings, with mixed risk of bias and performance metrics reported.
- **Overview of Tools:**
 - Identified 116 risk prediction tools across all included reviews.
 - Tools were divided based on the use of machine learning (ML) methods: 45% utilized ML (52/116) while 55% did not (64/116).
- **Data Collection and Model Development:**
 - Data used for model development varied: 19% used prospectively collected data (22/116), 40% used retrospectively collected data (46/116, including 18 ML-based models using electronic health records), and data collection methods were not reported for 41% (48/116).
 - Common settings for model development included hospital inpatients (34 tools) and long-term care settings such as rehabilitation units or nursing homes (20 tools).

3. Results (continued)

Tool characteristics	ML-based models (N=52, 45%)	Non-ML tools (N=64, 55%)	Total (N=116)
No. of included reviews ^a considered in			
0	0 (0)	13 (20)	13 (11)
1	30 (58)	23 (36)	53 (46)
2	8 (15)	9 (14)	17 (15)
>2	14 (27)	19 (30)	33 (28)
Development study details			
Median (range) year of publication	2020 (2000 – 2022)	1998 (1962 – 2015)	2006 (1962 – 2022)
Source of data			
Prospective	4 (8)	18 (28)	22 (19)
Retrospective	36 (69)	10 (16)	46 (40)
Setting	NS	36 (56)	48 (41)
Hospital	13 (25)	11 (17)	24 (21)
Long-term care (incl. end-of-life and rehab)	6 (11)	14 (22)	20 (17)
Acute care (incl. surgical and ICU)	28 (54)	24 (38)	52 (45)
Mixed settings	1 (2)	1 (2)	2 (2)
Other	2 (4)	2 (3)	4 (3)
NS	2 (4)	12 (19)	14 (12)
Study population age			
Adults	31 (60)	34 (53)	65 (56)
Any	5 (10)	3 (5)	8 (7)
Any	16 (31)	27 (42)	43 (37)
Baseline condition			
Pis at baseline	1 (2)	0 (0)	1 (1)
No Pis at baseline	11 (21)	19 (30)	30 (26)
NS	40 (77)	45 (70)	85 (73)
Development methods			
Development method/algorithms ^b			
ML algorithms	52 (100)	0 (0)	52 (45)
Logistic regression	35 (67)	15 (23) †	50 (43)
Cox regression	0 (0)	3 (5)	3 (3)
Clinical expertise	3 (6)	3 (5)	6 (5)
NS	0 (0)	44 (69) ‡	44 (38)
Internal validation method			
Cross-validation	21 (40)	3 (5) †*	24 (21)
Data splitting	11 (21)	0 (0)	11 (9)
Not done / NS	20 (39)	61 (95)	81 (69)
Median (range) no. of final predictive events	6 (1–33)	8 (2–15)	7 (2–23)
Study cohort			
Median (range) total sample size	3,000 (271 – 1,252,313)	320 (15 – 331,150)	686 (15 – 1,252,313)
Median (range) number of events	206 (8 – 86,410)	51 (9 – 1,350)	94 (8 – 86,410)
Median (range) proportion of events	10.65% (0.42% – 80.00%)	14.84% (1.18% – 46.67%)	14.40% (0.42% – 80.00%)

Table 3. Summary of tool characteristics, from review-level data

- **Model Characteristics and Internal Validation:**
 - Sample sizes, where reported ($n=92$), ranged dramatically from 1,577 to over 1.25 million.
 - Internal validation approaches were poorly documented, with specific methods unidentified for 70% of models (81/116).
- **Predictors Used in Models:**
 - Information on included predictors was detailed for 53 of the 116 tools.
 - Most common predictors included mobility (51%), predisposing diseases/conditions (49%), medication treatment/care (42%), and continence (42%).
 - Other frequently mentioned predictors were age, nutrition, mental status, activity, skin conditions, and lab values.
 - Some tools also integrated established risk prediction scales such as Braden scores (six tools) and the Norton score (one tool).

Predictor category	No. of tools predictor appears in
Mobility	27 (51)
Pre-disposing conditions	26 (49)
Contra-indications	22 (42)
Receiving medical treatment/care	22 (42)
Age	21 (40)
Mental Status	20 (38)
Nutrition	20 (38)
Activity	19 (36)
Laboratory values	18 (34)
Skin	18 (34)
General Health	18 (30)
Body	13 (25)
Gender	11 (21)
Surgery duration	7 (13)
Ability to ambulate	6 (11)
Braided score	6 (11)
Pressure injury	6 (11)
Friction, shear, pressure	5 (9)
Medical unit, ward, visit	5 (9)
Length of stay	4 (8)
Hygiene	3 (6)
Ethnicity	2 (4)
Pain	2 (4)
Smoking	2 (4)
'Special' (not explained)	2 (4)
Isolation	1 (2)
Norton score	1 (2)

Table 4. Summary of tool characteristics, from review-level data
Figures are given as count (% out of 53 tools with information on predictors). Note that multiple predictors may fall within the same predictor category. For instance, category 'skin' may encompass both 'skin moisture' and 'skin integrity', with the frequency count reflecting the entire predictor category rather than individual predictors

- **Limitations**
- **Review Standards and Search Strategy:**
 - Conducted following Cochrane guidelines with a highly sensitive search designed by an experienced information specialist.
 - Excluded non-English publications due to time and resources but used them to identify additional models where possible.
- **Quality Assessment Challenges:**
 - Used AMSTAR-2 for quality assessment, which is not specifically designed for diagnostic or prognostic studies.
 - Adaptations were made to AMSTAR-2, but these focused more on reporting quality rather than methodological quality.
 - Potential need for establishing specific criteria for assessing systematic reviews of prediction models.
- **Limitations in Review Findings:**
 - Significant gaps in detail about risk prediction models and their performance as reported by the systematic reviews.
 - Reporting quality likely affected by the lack of contemporary guidelines during the development of many traditional (non-ML) risk prediction tools, most of which predate 2000. Contrastingly, ML-based model published mostly post-2000 (median year 2020), benefitted from more recent reporting guidelines.

4. CONCLUSIONS

- **Evidence Scope and Review Quality:**
 - Extensive evidence on risk prediction scales, tools, and models for pressure injuries (PIs) summarized across multiple systematic reviews of varied methodological quality.
 - Only five systematic reviews comprehensively reported on the development and validation of models specifically designed to predict the risk of PIs.
- **Current Standards and Model Validation:**
 - Many available models, including those utilizing machine learning (ML), fail to meet current standards for development and reporting of risk prediction models.
 - Most models have not undergone validation outside the original populations in which they were developed, limiting their generalizability.
- **Optimal Model Identification and Research Needs:**
 - Identifying the optimal risk prediction model for PI requires a high-quality systematic review focused on primary literature adhering to stringent methodological standards.
 - Current lack of consensus on the best risk prediction model for PI underscores the need for more standardized and rigorous research methodologies in future studies.

This overview emphasizes the critical need for enhanced methodological rigor and standardized reporting in the development and evaluation of risk prediction models to improve their reliability and applicability in clinical settings.

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