Review article



The roles of tertiary lymphoid structures in chronic diseases

In the format provided by the authors and unedited

Supplementary table 1 | Frequency of TLS-positive cases in cohorts of different tumours extracted from studies reporting histological TLS assessment

| Included in organ comparison analysis | Cancer subtype | Organ | Disease details | Treatment before surgery | total cases analysed | # of TLS+ cases | % of TLS- positive cases | Details on TLS assessment method and region | Reference |
|---------------------------------------|-------------------|----------|--|-----------------------------|-------------------------|--------------------|--------------------------------|---|---|
| Yes | IBC | Breast | Invasive ductal carcinoma, mixed subtypes | untreated | 134 | 17 | 12.69 | whole slide H&E, whole tissue area | 1 |
| Yes | DCIS | Breast | Ductal carcinoma in situ | untreated | 68 | 5 | 7.35 | whole slide H&E, whole tissue area | 1 |
| Yes | HNSCC | Tongue | Oral tongue squamous cell carcinoma | untreated | 310 | 263 | 84.84 | whole slide H&E, peritumoral area | 2 |
| Yes | HCC | Liver | Hepatocellular carcinoma | untreated | 273 | 129 | 47.25 | whole slide H&E, whole tissue area, TLS assessed intratumorally | 3 |
| Yes | PDAC | Pancreas | Pancreatic ductal adenocarcinoma | untreated | 104 | 101 | 97.12 | whole slide IHC for CD20, tumor-stroma interface | 4 |
| Yes | TNBC | Breast | Metaplastic breast cancer (triple negative breast cancer-like) | untreated | 60 | 36 | 60.00 | whole slide H&E, whole tissue area, number of positive cases extrapolated from survival graph | 5 |
| Yes | NSCLC | Lung | Non-small cell lung cancer, early stage | untreated | 74 | 64 | 86.49 | whole slide H&E and IHC for DC-LAMP and CD3, whole tissue area | 6 |
| Yes | BC_HER2- | Breast | HER2- invasive breast cancer and ductal carcinoma in situ | untreated | 210 | 69 | 32.86 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | BC_HER2+ | Breast | HER2+ invasive breast cancer and ductal carcinoma in situ | untreated | 40 | 26 | 65.00 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | BC_HER2- | Breast | ER- invasive breast cancer and ductal carcinoma in situ | untreated | 41 | 31 | 75.61 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | BC_HER2- | Breast | PR- invasive breast cancer and ductal carcinoma in situ | untreated | 56 | 36 | 64.29 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | BC_HER2+ | Breast | ER+ invasive breast cancer and ductal carcinoma in situ | untreated | 208 | 64 | 30.77 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | BC_HER2+ | Breast | PR+ invasive breast cancer and ductal carcinoma in situ | untreated | 120 | 38 | 31.67 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| Yes | HNSCC | Larynx | laryngeal squamous cell carcinoma | untreated | 24 | 5 | 20.83 | whole slide H&E, whole tissue area | 8 |
| Yes | NSCLC | Lung | Non-small cell lung cancer | untreated | 147 | 86 | 58.50 | whole slide IHC for PNAD next to dense lymphocytic clusters, 10 40x regions per slide | 9 |
| Yes | HCC | Liver | Hepatocellular carcinoma | untreated | 112 | NA | 33.00 | whole slide multiplexIF, whole tissue area | 10 |
| Yes | NSCLC | Lung | Non-small cell lung cancer | untreated | 103 | 80 | 77.67 | whole slide H&E, whole tissue area | 11 |
| Yes | PDAC_TMBhi | Pancreas | Pancreatic ductal adenocarcinoma, SHM high (>10) | untreated | 17 | 14 | 82.35 | whole slide IHC for CD20 and CD3, multiplex IF, whole tissue area | 12 |
| Yes | PDAC_TMBlo | Pancreas | Pancreatic ductal adenocarcinoma, SHM low (<=10) | untreated | 42 | 5 | 11.90 | whole slide IHC for CD20 and CD3, multiplex IF, whole tissue area | 12 |
| Yes | PDAC | Pancreas | Pancreatic ductal adenocarcinoma | untreated | 174 | 44 | 25.29 | whole slide H&E, whole tissue area | 12 |
| Yes | BC_HER2- | Breast | HR- invasive breast cancer | untreated | 41 | 19 | 46.34 | whole slide IHC for CD20, whole tissue area | 10.3969/j.issn.167 3-4254.2013.05.25 |
| Yes | BC_HER2+ | Breast | HR+ invasive breast cancer | untreated | 89 | 30 | 33.71 | whole slide IHC for CD20, whole tissue area | 10.3969/j.issn.167 3-4254.2013.05.25 |
| Yes | GACA | Stomach | Gastric adenocarcinoma, intestinal type | untreated | 6 | 6 | 100.00 | whole slide IHC for CD20, whole tissue area | 13 |
| Yes | PDAC | Pancreas | Pancreatic ductal adenocarcinoma | untreated | 308 | 308 | 100.00 | whole slide H&E, whole tissue area | 14 |

| Yes | TNBC | Breast | Triple negative breast cancer | untreated | 42 | 21 | 50.00 | whole slide IHC for CD21, CD20, whole tissue area | 15 |
|-----|------------|-------------------|---|-----------|-----|-----|--------|--|----|
| Yes | NSCLC | Lung | Lung adenocarcinoma | untreated | 146 | 57 | 39.04 | whole slide H&E, whole tissue area | 16 |
| Yes | MIBC | Bladder | Muscle invasive bladder cancer | untreated | 16 | 13 | 81.25 | whole slide multiplex IHC, whole tissue area | 17 |
| Yes | NonMIBC | Bladder | Non- Muscle invasive bladder cancer | untreated | 12 | 9 | 75.00 | whole slide multiplex IHC, whole tissue area | 17 |
| Yes | OVCA | Ovary and | High-grade serous ovarian cancer | untreated | 30 | 17 | 56.67 | whole slide multiplex IHC, whole tissue area | 18 |
| Yes | TNBC | omentum Breast | Triple negative breast cancer | untreated | 769 | 733 | 95.32 | whole slide H&E, whole tissue area | 19 |
| Yes | BC_HER2+ | Breast | HER2+ invasive breast cancer | untreated | 447 | 402 | 89.93 | whole slide H&E, whole tissue area, TLS found peritumorally | 20 |
| Yes | BC_HER2- | Breast | HR-HER2+ invasive breast cancer | untreated | 240 | 186 | 77.50 | whole slide H&E, whole tissue area, TLS found peritumorally | 20 |
| Yes | BC_HR+ | Breast | HR+HER2+ invasive breast cancer | untreated | 207 | 116 | 56.04 | whole slide H&E, whole tissue area, TLS found peritumorally | 20 |
| Yes | BC_HER2+ | Breast | HER2+ invasive breast cancer | untreated | 168 | 65 | 38.69 | whole slide H&E, whole tissue area | 21 |
| Yes | BC_HER2- | Breast | HR-HER2+ invasive breast cancer | untreated | 78 | NA | 43.59 | whole slide H&E, whole tissue area | 21 |
| Yes | BC_HR+ | Breast | HR+HER2+ invasive breast cancer | untreated | 90 | NA | 34.44 | whole slide H&E, whole tissue area | 21 |
| Yes | BC_HER2- | Breast | HER2- invasive breast cancer | untreated | 191 | 59 | 30.89 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 22 |
| Yes | BC_HER2+ | Breast | HER2+ invasive breast cancer | untreated | 54 | 33 | 61.11 | whole slide H&E and IHC for CD20, CD3, CD23, | 22 |
| Yes | BC_HR- | Breast | ER- invasive breast cancer | untreated | 89 | 46 | 51.69 | whole tissue area whole slide H&E and IHC for CD20, CD3, CD23, | 22 |
| Yes | BC_HR- | Breast | PR- invasive breast cancer | untreated | 95 | 50 | 52.63 | whole tissue area whole slide H&E and IHC for CD20, CD3, CD23, | 22 |
| Yes | BC_HR+ | Breast | ER+ invasive breast cancer | untreated | 157 | 46 | 29.30 | whole tissue area whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 22 |
| Yes | BC_HR+ | Breast | PR+ invasive breast cancer | untreated | 150 | 42 | 28.00 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 22 |
| Yes | TNBC | Breast | Triple negative breast cancer | untreated | 45 | 22 | 48.89 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 22 |
| Yes | MIBC | Bladder | Urinary bladder | untreated | 51 | 28 | 54.90 | 3mm TMA cores with TC and IM by IF, sequential IHC | 23 |
| Yes | RCC | Kidney | Clear cell renal cell carcinoma | untreated | 105 | 33 | 31.43 | | 23 |
| Yes | CRC | Liver | Metastatic colorectal cancer | untreated | 14 | 12 | 85.71 | whole slide IHC for CD20, whole tissue area, TLS present peritumorally | 24 |
| Yes | CRC | Colon | Colorectal cancer | untreated | 843 | 230 | 27.28 | whole slide H&E, whole tissue area, TLS present peritumorally | 25 |
| Yes | MIBC_TMBhi | Bladder | Muscle invasive bladder cancer, mutation high | untreated | 151 | 117 | 77.48 | whole slide H&E, whole tissue area | 26 |
| Yes | MIBC_TMBlo | Bladder | Muscle invasive bladder cancer, mutation low | untreated | 153 | 107 | 69.93 | whole slide H&E, whole tissue area | 26 |
| Yes | MIBC | Bladder | Muscle invasive bladder cancer | untreated | 40 | 40 | 100.00 | whole slide multiplexIF, whole tissue area | 26 |
| Yes | CRC_TMBhi | Colon | Colorectal cancer, microsatellite instable | untreated | 15 | 15 | 100.00 | whole slide multiplexIF, peritumoral area | 27 |
| Yes | CRC_TMBlo | Colon | Colorectal cancer, microsatellite stable | untreated | 94 | 91 | 96.81 | whole slide multiplexIF, peritumoral area | 27 |
| | | | | | | | | | |

| Yes | CRC_TMBhi | Colon | Colorectal cancer, microsatellite instable | untreated | 189 | 111 | 58.73 | whole slide H&E, whole tissue area | 28 |
|-----|------------|--|--|----------------------------|------|-----|--------|---|----|
| Yes | CRC_TMBlo | Colon | Colorectal cancer, microsatellite stable | untreated | 1299 | 588 | 45.27 | whole slide H&E, whole tissue area | 28 |
| Yes | HNSCC_HPV- | Tongue | HPV- head and neck squamous cell carcinoma | untreated | 13 | 13 | 100.00 | whole slide IHC for CD20, whole tissue area | 29 |
| Yes | HNSCC_HPV- | Tonsil | HPV- head and neck squamous cell carcinoma | untreated | 18 | 18 | 100.00 | whole slide IHC for CD20, whole tissue area | 29 |
| Yes | HNSCC_HPV+ | Tongue | HPV+ head and neck squamous cell carcinoma | untreated | 15 | 12 | 80.00 | whole slide IHC for CD20, whole tissue area | 29 |
| Yes | HNSCC_HPV+ | Tonsil | HPV+ head and neck squamous cell carcinoma | untreated | 10 | 9 | 90.00 | whole slide IHC for CD20, whole tissue area | 29 |
| Yes | NSCLC | Lung | Lung squamous cell carcinoma | untreated | 69 | 68 | 98.55 | whole slide H&E, peritumoral area | 30 |
| Yes | BC_HER2+ | Breast | HER2+ invasive ductal cancer | untreated | 48 | 48 | 100.00 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 31 |
| Yes | TNBC | Breast | Triple negative breast cancer | untreated | 20 | 18 | 90.00 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 31 |
| Yes | OVCA | Ovary and omentum | High-grade serous ovarian cancer | untreated | 147 | 14 | 9.52 | whole slide IHC for CD20 and DC-LAMP, whole tissue area | 32 |
| Yes | OVCA | Ovary and omentum | High-grade serous ovarian cancer | untreated | 62 | 39 | 62.90 | whole slide IHC for CD20, whole tissue area | 33 |
| Yes | OVCA | Ovary and omentum | High-grade serous ovarian cancer | untreated | 35 | 22 | 62.86 | whole slide IHC for CD20, whole tissue area | 33 |
| Yes | MIBC | Bladder | Muscle invasive bladder cancer | untreated | 31 | 25 | 80.65 | whole slide multiplex IF, whole tissue area, most TLS peritumorally | 34 |
| Yes | CRC | Colon | Colorectal cancer | untreated | 418 | 411 | 98.33 | whole slide H&E, whole tissue area, TLS present peritumorally | 35 |
| Yes | HNSCC_HPV- | Oral cavity excluding base of tongue and tonsils | HPV- oral squamous cell carcinoma | untreated | 68 | 16 | 23.53 | whole slide IHC for CD20, CD21, BCL6, whole tissue area | 36 |
| Yes | HNSCC_HPV+ | Oral cavity excluding base of tongue and tonsils | HPV+ oral squamous cell carcinoma | untreated | 6 | 1 | 16.67 | whole slide IHC for CD20, CD21, BCL6, whole tissue area | 36 |
| Yes | CRC | Colon | Colorectal cancer | untreated | 67 | 58 | 86.57 | multiplex IF, whole tissue area, TLS present peritumorally | 37 |
| Yes | NSCLC | Lung | Lung adenocarcinoma, without ALK rearrangement and driver gene mutations | untreated | 9 | 8 | 88.89 | whole slide IHC for CD20, whole tissue area | 38 |
| Yes | NSCLC | Lung | Lung adenocarcinoma, with ALK rearrangement | untreated | 39 | 30 | 76.92 | whole slide IHC for CD20, whole tissue area | 38 |
| Yes | HCC | Liver | Hepatocellular carcinoma | untreated | 170 | 142 | 83.53 | whole slide H&E, peritumoral area | 39 |
| No | | Breast | Invasive ductal carcinoma, mixed subtypes | Chemotherapy | 35 | 0 | 0.00 | whole slide H&E, whole tissue area | 1 |
| No | | Breast | Invasive ductal carcinoma, mixed subtypes | untreated+che motherapy | 312 | 37 | 11.86 | whole slide H&E, whole tissue area | 1 |
| No | | Prostate | Prostate adenocarcinoma | untreated | 17 | NA | NA | whole slide H&E, whole tissue area | 40 |
| No | | Breast | invasive breast cancer and ductal carcinoma in situ | untreated | 290 | 146 | 50.34 | whole slide H&E and IHC for CD20, CD3, CD21, BCL6, PNAD, whole tissue area | 7 |
| No | | Lung | Non-small cell lung cancer, advanced stage | Chemotherapy | 122 | NA | NA | whole slide IHC for DC-LAMP and CD20, whole tissue area | 41 |
| No | | Lung | Non-small cell lung cancer, early stage | untreated | 74 | NA | NA | whole slide IHC for DC-LAMP and CD20, whole tissue area | 41 |

| No | Lung | Non-small cell lung cancer without underlying COPD | untreated | 376 | NA | NA | whole slide IHC for DC-LAMP and CD8, whole tissue area | 42 |
|----|----------|--|----------------------------|------|-----|-------|--|---|
| No | Lung | Non-small cell lung cancer | NA | 91 | NA | NA | whole slide H&E, whole tissue area | 43 |
| No | Breast | Invasive breast cancer, mixed subtypes | untreated | 70 | NA | NA | whole slide IHC, Tfh used as surrogate for TLS, whole tissue area | 44 |
| No | Pancreas | Pancreatic ductal adenocarcinoma, SHM high + low | untreated | 63 | 29 | 46.03 | whole slide IHC for CD20 and CD3, multiplex IF, whole tissue area | 12 |
| No | Breast | Invasive breast cancer, mixed subtypes | untreated | 130 | 49 | 37.69 | whole slide IHC for CD20, whole tissue area | 10.3969/j.issn.167 3-4254.2013.05.25 |
| No | Stomach | Gastric adenocarcinoma | untreated | 1033 | NA | NA | whole slide H&E and IHC for CD20, CD21, whole tissue area | 45 |
| No | Stomach | Gastric adenocarcinoma | Chemotherapy | 42 | NA | NA | whole slide IHC for CD20, whole tissue area | 46 |
| No | Stomach | Gastric adenocarcinoma | untreated | 40 | NA | NA | whole slide IHC for CD20, whole tissue area | 46 |
| No | Stomach | Gastritits with intestinal metaplasia, non-tumor | untreated | 10 | 0 | 0.00 | whole slide IHC for CD20, whole tissue area | 13 |
| No | Brain | Metastatic breast cancer | untreated+che motherapy | 37 | 0 | 0.00 | whole slide H&E, whole tissue area | 47 |
| No | Breast | Metastatic breast cancer, primary tumor | untreated+che motherapy | 245 | 112 | 45.71 | whole slide H&E, whole tissue area | 47 |
| No | Liver | Metastatic breast cancer | untreated+che motherapy | 84 | 13 | 15.48 | whole slide H&E, whole tissue area | 47 |
| No | Lung | Metastatic breast cancer | untreated+che motherapy | 127 | 48 | 37.80 | whole slide H&E, whole tissue area | 47 |
| No | Ovary | Metastatic breast cancer | untreated+che motherapy | 25 | 0 | 0.00 | whole slide H&E, whole tissue area | 47 |
| No | Liver | Hepatocellular carcinoma | untreated | 303 | 102 | 33.66 | TMA H&E, intratumoral area | 48 |
| No | Liver | Hepatocellular carcinoma | untreated | 159 | 54 | 33.96 | TMA H&E, intratumoral area | 48 |
| No | Liver | Hepatic hemangioma, non-tumor | untreated | 50 | 4 | 8.00 | TMA H&E, whole tissue | 48 |
| No | Breast | Invasive breast cancer, mixed subtypes | untreated | 255 | 101 | 39.61 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 22 |
| No | Rectum | Rectal cancer | untreated | 60 | NA | NA | whole slide IHC of CD3, CD83, whole tissue area, TLS peritumorally | 49 |
| No | Liver | Metastatic colorectal cancer | Chemotherapy | 51 | 33 | 64.71 | whole slide IHC for CD20, whole tissue area, TLS present peritumorally | 24 |
| No | Liver | Metastatic colorectal cancer | untreated+che motherapy | 65 | 45 | 69.23 | whole slide IHC for CD20, whole tissue area, TLS present peritumorally | 24 |
| No | Omentum | High-grade serous ovarian cancer | Chemotherapy | 31 | NA | NA | whole slide IHC for CD20, whole tissue area | 50 |
| No | Omentum | High-grade serous ovarian cancer | untreated | 10 | NA | NA | whole slide IHC for CD20, whole tissue area | 50 |
| No | Liver | Hepatocellular carcinoma | untreated | 145 | 59 | 40.69 | whole slide IHC for CD3, CD8, CD20, CD208, CD21, intratumoral area | 51 |
| No | Liver | Hepatocellular carcinoma | untreated | 120 | 48 | 40.00 | whole slide IHC for CD3, CD8, CD20, CD208, CD21, intratumoral area | 51 |
| No | Bladder | Muscle invasive bladder cancer | untreated | 300 | 58 | 19.33 | whole slide H&E, intratumoral area | 26 |
| No | Bladder | Muscle invasive bladder cancer, mutation low+high | untreated | 304 | 223 | 73.36 | whole slide H&E, whole tissue area | 26 |
| No | Bladder | Muscle invasive bladder cancer | Chemotherapy | 30 | 29 | 96.67 | whole slide multiplexIF, whole tissue area | 26 |

| No | Colon | Colorectal cancer, microsatellite stable and unstable | untreated | 109 | 106 | 97.25 | whole slide multiplexIF, peritumoral area | 27 |
|----|--|---|--|------|-----|--------|---|----|
| No | Lung | Non-small cell lung cancer, advanced stage | Chemotherapy | 122 | NA | NA | whole slide IHC for DC-LAMP and CD8, whole tissue area | 52 |
| No | Lung | Non-small cell lung cancer, advanced stage | untreated | 39 | NA | NA | whole slide IHC for DC-LAMP and CD8, whole tissue area | 52 |
| No | Lung | Metastatic colorectal cancer | untreated | 140 | NA | NA | whole slide IHC for DC-LAMP and CD8, whole tissue area | 52 |
| No | Lung | Metastatic renal cell carcinoma | untreated | 52 | NA | NA | whole slide IHC for DC-LAMP and CD8, whole tissue area | 52 |
| No | Colon | Colorectal cancer, microsatellite stable and unstable | untreated | 1663 | 784 | 47.14 | whole slide H&E, whole tissue area | 28 |
| No | Tonsil and base of Tongue | HPV- head and neck squamous cell carcinoma | untreated | 25 | 25 | 100.00 | whole slide IHC for CD20, whole tissue area | 29 |
| No | Tonsil and base of Tongue | HPV+ head and neck squamous cell carcinoma | untreated | 25 | 21 | 84.00 | whole slide IHC for CD20, whole tissue area | 29 |
| No | Lung | Metastatic colorectal cancer | untreated+che motherapy | 57 | 45 | 78.95 | whole slide IHC of CD8, FOXP3 whole tissue area | 53 |
| No | Lung | Lung squamous cell carcinoma | Chemotherapy | 51 | 47 | 92.16 | whole slide H&E, peritumoral area | 30 |
| No | Lung | Lung squamous cell carcinoma | tumor non- related corticosteroids | 18 | 18 | 100.00 | whole slide H&E, peritumoral area | 30 |
| No | Lung | Lung squamous cell carcinoma | untreated+che motherapy | 138 | 133 | 96.38 | whole slide H&E, whole tissue area | 30 |
| No | Breast | Invasive ductal carcinoma, mixed subtypes | untreated | 167 | 148 | 88.62 | whole slide H&E and IHC for CD20, CD3, CD23, whole tissue area | 31 |
| No | Breast | Triple negative breast cancer | Chemotherapy | 108 | 93 | 86.11 | whole slide H&E, whole tissue area | 54 |
| No | Lung | Non-small cell lung cancer with underlying COPD | Brochodilators | 90 | NA | NA | whole slide IHC for CD20 and CD3, whole tissue area | 55 |
| No | Lung | Non-small cell lung cancer without underlying COPD | untreated | 43 | NA | NA | whole slide IHC for CD20 and CD3, whole tissue area | 55 |
| No | Ovary and omentum | High-grade serous ovarian cancer | untreated | 97 | 61 | 62.89 | whole slide IHC for CD20, whole tissue area | 33 |
| No | Colon | Colorectal cancer | Chemotherapy or RT | 32 | NA | NA | whole slide H&E, whole tissue area, TLS present peritumorally | 35 |
| No | Colon | Colorectal cancer | untreated | 117 | NA | NA | whole slide H&E, whole tissue area, TLS present peritumorally | 35 |
| No | Colon | Colorectal cancer | untreated+che motherapy+RT | 149 | 147 | 98.66 | whole slide H&E, whole tissue area, TLS present peritumorally | 35 |
| No | Liver | Hepatocellular carcinoma | untreated | 163 | 10 | 6.13 | whole slide H&E, whole tissue area, TLS assessed intratumorally | 56 |
| No | Oral cavity excluding base of tongue and tonsils | Oral squamous cell carcinoma, mixed HPV | untreated | 80 | 17 | 21.25 | whole slide IHC for CD20, CD21, BCL6, whole tissue area | 36 |
| No | Lung | Lung adenocarcinoma, mixed ALK status | untreated | 48 | 38 | 79.17 | whole slide IHC for CD20, whole tissue area | 38 |
| No | Liver | Hepatocellular carcinoma | untreated | 170 | 73 | 42.94 | whole slide H&E, intratumoral area | 39 |

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Supplementary table 2 | TLS in human kidney diseases

| Definition of TLS | Frequency (TLS positive patients/total patients) | Clinical findings | Ref |
|--|---|---|----------|
| IgA nephropathy* | / | 1 | l . |
| Clusters containing CD4/CD8/CD20/DC- SIGN ⁺ cells | 38.8% (59/152) | TLS ⁺ patients had higher sCr, proteinuria, BP and more severe glomerulonephritis than TLS ⁻ patients | 1 |
| T and B cell aggregates | 31.4% (328/1044) | TLS frequency correlated with risk of 50% decline in eGFR or kidney failure in patients with eGFR<60 ml/min/1.73 m ² ; TLS predicted IgAN progression in earlier stage disease | 2 |
| Membranous nephro | pathy* | · • | • |
| T and B cell aggregates | 34.2% (151/442) | TLS+ patients had higher sCr, proteinuria and BP, lower serum albumin, and lower remission rate than TLS-patients; TLSs were associated with anti-PLA2R antibody; TLS+ patients were older and frequently male | 3 |
| Lupus nephritis* | | | |
| T and B cell aggregates T and B cell | 31.3% (10/32) 12.5% (4/32) | In a few patients, follow-up biopsy samples showed mature TLSs despite immunosuppressive treatment | 4 |
| aggregates with FDCs | , , | | |
| T and B cell aggregates | 51.4% (35/68) | The presence of TLS was associated with clonal expansion, somatic hypermutation and immune complex | 5 |
| T and B cell aggregates with germinal centre | 5.9% (4/68) | deposition in tubular basement membranes | |
| ANCA-associated glo | | | T |
| T and B cell aggregates | 31.3% (5/16) | No clinical findings were reported. | 4 |
| T and B cell aggregates with FDCs | 6.3% (1/16) | | |
| T and B cell aggregates | 47.4% (53/112) | The presence of TLS was associated with higher risk of AKI requiring dialysis and kidney failure | 6 |
| T and B cell aggregates with nodular B cell aggregates | 13.4% (15/112) | | |
| Acute kidney allograf | t rejection* | | |
| B cell clusters | 45.0% (9/20) | B cell density positively correlated with glucocorticoid resistance and poor graft outcomes | 7 |
| B cell clusters | 59.2% (71/120) | The presence of TLSs was not associated with time to ACR, steroid resistance, sCr or graft loss | 8 |
| Clinically stable kidne | L ev allografts* | 1 | <u> </u> |
| T and B cell aggregates with signs of proliferation | 0 months: 3.8% (8/210); 1 month: 46.9% (90/192); 6 months: 53.4% | The presence of TLS with FDCs in allografts at 1-year biopsy was associated with higher risk of kidney function decline after transplantation compared with allografts without TLS or those with TLSs but no FDCs | 9 |

| | (87/163) 12 months: | | |
|------------------------|----------------------------------|-------------------------------------|----|
| | 53.4% (108/185) | | |
| T and B cell | 0 months: 1.4% | | |
| aggregates with signs | (3/210); 1 month: | | |
| of proliferation and | 3.6% (7/192); 6 | | |
| FDCs | months: 8.6% | | |
| | (14/163); 12 months: | | |
| | 18.9% (35/185) | | |
| Kidney allografts witl | h chronic rejection [‡] | | |
| B cell clusters with | 95% (19/20) | No clinical findings were reported. | 10 |
| FDCs | | | |
| Complicated pyelone | ohritic kidneys [‡] | | |
| T and B cell | 100.0% (16/16) | No clinical findings were reported. | 11 |
| aggregates with sign | | | |
| of proliferation | | | |

*Needle biopsy samples; *Surgically resected kidneys. sCr, serum creatinine; BP, blood pressure; PLA2R, phospholipase A2 receptor; FDCs, follicular dendritic cells; ANCA, Antineutrophil Cytoplasmic Antibody; AKI, acute kidney injury; ACR, acute cellular rejection; BCR, B cell receptor; CKD, chronic kidney disease.

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